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1                                   **Cancer Treatment**

2

3       **Field of the Invention**

4

5       The present invention relates to cancer treatment.  
6       In particular, it relates to methods and  
7       compositions for the treatment of cancer, including  
8       cancers characterised by p53 mutations..  
9

10       **Background to the Invention**

11

12       5-FU<sup>4</sup> is widely used in the treatment of a range of  
13       cancers including colorectal, breast and cancers of  
14       the aerodigestive tract. The mechanism of cytotoxicity  
15       of 5-FU has been ascribed to the misincorporation of  
16       fluoronucleotides into RNA and DNA and to the  
17       inhibition of the nucleotide synthetic enzyme  
18       thymidylate synthase (TS) (Longley et al., 2003). TS  
19       catalyses the conversion of deoxyuridine monophosphate  
20       (dUMP) to deoxythymidine monophosphate (dTMP) with  
21       5,10-methylene tetrahydrofolate (CH<sub>2</sub>THF) as the methyl  
22       donor. This reaction provides the sole intracellular

1 source of thymidylate, which is essential for DNA  
2 synthesis and repair. The 5-FU metabolite  
3 fluorodeoxyuridine monophosphate (FdUMP) forms a  
4 stable complex with TS and CH<sub>2</sub>THF resulting in enzyme  
5 inhibition (Longley et al., 2003). Recently, more  
6 specific folate-based inhibitors of TS have been  
7 developed such as tomudex (TDX) and Alimta (MTA),  
8 which form a stable complex with TS and dUMP that  
9 inhibits binding of CH<sub>2</sub>THF to the enzyme (Hughes et  
10 al., 1999; Shih et al., 1997). TS inhibition causes  
11 nucleotide pool imbalances that result in S phase cell  
12 cycle arrest and apoptosis (Aherne et al., 1996;  
13 Longley et al., 2002; Longley et al., 2001).  
14 Oxaliplatin is a third generation platinum-based DNA  
15 damaging agent that is used in combination with 5-FU  
16 in the treatment of advanced colorectal cancer  
17 (Giacchetti et al., 2000). Drug resistance is a major  
18 factor limiting the effectiveness of chemotherapies.  
19 The topoisomerase-1 inhibitor irinotecan (CPT-11) and  
20 the DNA damaging agent oxaliplatin are now being used  
21 in conjunction with 5-FU for the treatment of  
22 metastatic colorectal cancer, having demonstrated  
23 improved response rates compared to treatment with 5-  
24 FU alone (40-50% compared to 10-15%) (10, 11). Despite  
25 these improvements, the vast majority of responding  
26 patients relapse, with median survival times of only  
27 22-24 months. Clearly, new approaches are needed for  
28 the treatment of this disease.

29

30 Death receptors such as Fas and the TRAIL (tumour  
31 necrosis factor (TNF)-related apoptosis-inducing  
32 ligand) receptors DR4 (TRAIL-R1) and DR5 (TRAIL-R2)

1 trigger death signals when bound by their natural  
2 ligands (1,2). Ligand binding to the death receptors  
3 leads to recruitment of the adaptor protein FADD  
4 (Fas-associated death domain), which in turn  
5 recruits procaspase 8 zymogens to form the death-  
6 inducing signalling complex (DISC) (Nagata, 1999).  
7 Procaspase 8 molecules become activated at the DISC  
8 and subsequently activate pro-apoptotic downstream  
9 molecules such as caspase 3 and BID. FasL expression  
10 is up-regulated in most colon tumours, and it has  
11 been postulated that tumour FasL induces apoptosis  
12 of Fas-sensitive immune effector cells (O'Connell et  
13 al., 1999). This mechanism of immune escape requires  
14 that tumour cells develop resistance to Fas-mediated  
15 apoptosis to prevent autocrine and paracrine tumour  
16 cell death.

17  
18 A key inhibitor of Fas signaling is c-FLIP, which  
19 inhibits procaspase 8 recruitment and processing at  
20 the DISC (Krueger et al., 2001). Differential  
21 splicing gives rise to long (c-FLIP<sub>L</sub>) and short (c-  
22 FLIP<sub>S</sub>) forms of c-FLIP, both of which bind to FADD  
23 within the DISC. c-FLIP<sub>S</sub> directly inhibits caspase 8  
24 activation at the DISC, whereas c-FLIP<sub>L</sub> is first  
25 cleaved to a p43 truncated form that inhibits  
26 complete processing of procaspase 8 to its active  
27 subunits. c-FLIP also inhibits procaspase 8  
28 activation at DISCs formed by the TRAIL (TNF-related  
29 apoptosis-inducing ligand) death receptors DR4  
30 (TRAIL-R1) and DR5 (TRAIL-R2) (Krueger et al.,  
31 2001). In addition to blocking caspase 8 activation,  
32 DISC-bound c-FLIP has been reported to promote

1 activation of the ERK, PI3-kinase/Akt and NF- $\kappa$ B  
2 signaling pathways (Krueger et al., 2001). Thus, c-  
3 FLIP potentially converts death receptor signaling  
4 from pro- to anti-apoptotic by activating intrinsic  
5 survival pathways. Significantly, c-FLIP<sub>L</sub> has been  
6 found to be overexpressed in colonic adenocarcinomas  
7 compared to matched normal tissue, suggesting that  
8 c-FLIP may contribute to in vivo tumour  
9 transformation (Ryu et al., 2001).

10

#### 11 **Summary of the Invention**

12

13 As described herein and, as shown in our co-pending  
14 PCT application filed on the same day as the present  
15 application and claiming priority from GB patent  
16 application 0327493.3, the present inventors have  
17 shown that by combining treatment using a death  
18 receptor ligand, such as an anti FAS antibody, for  
19 example, CH-11, with a chemotherapeutic agent such  
20 as 5-FU or an antifolate drug, such as raltitrexed  
21 (RTX) or pemetrexed (MTA, Alimta), a synergistic  
22 effect is achieved in the killing of cancer cells.  
23 However, the synergistic effect achieved was  
24 abrogated in cancer cells which overexpress c-FLIP.

25

26 As described in the Examples, in cell lines which  
27 demonstrated overexpression of c-FLIP and associated  
28 resistance to chemotherapy e.g 5-FU induced  
29 apoptosis, inhibition of FLIP expression reversed  
30 the resistance to chemotherapy-induced apoptosis.  
31 On further investigating this effect, the inventors

1 tested a number of cell lines having a p53 mutation  
2 or p53 null genotype.

3  
4 To their surprise, the inventors observed that down-  
5 regulation of c-FLIP markedly enhanced apoptosis in  
6 response to certain chemotherapeutic agents in the  
7 p53 mutant cells, which are usually highly resistant  
8 to the particular chemotherapeutic agents. This  
9 surprising observation enables the use of  
10 combinations of such cFLIP inhibitors and  
11 chemotherapeutic agents in the treatment of cancers  
12 associated with p53 mutations.

13  
14 Accordingly, in a first aspect of the present  
15 invention, there is provided a method of killing  
16 cancer cells having a p53 mutation, comprising  
17 administration to said cells of:

- 18 (a) a c-FLIP inhibitor and  
19 (b) a chemotherapeutic agent, wherein the  
20 chemotherapeutic agent is a thymidylate synthase  
21 inhibitor, a platinum cytotoxic agent or a  
22 topoisomerase inhibitor.

23  
24 In a second aspect, there is provided a method of  
25 treating cancer associated with a p53 mutation  
26 comprising administration to a subject in need  
27 thereof of

- 28 (a) a c-FLIP inhibitor and  
29 (b) a chemotherapeutic agent, wherein the  
30 chemotherapeutic agent is a thymidylate synthase  
31 inhibitor, a platinum cytotoxic agent or a

1     topoisomerase inhibitor.

2

3     A third aspect of the invention comprises the use of

4     (a) a c-FLIP inhibitor and

5     (b) a chemotherapeutic agent, wherein the

6     chemotherapeutic agent is a thymidylate synthase

7     inhibitor, a platinum cytotoxic agent or a

8     topoisomerase inhibitor

9     in the preparation of a medicament for treating

10    cancer associated with a p53 mutation.

11

12    A fourth aspect provides a pharmaceutical

13    composition for the treatment of a cancer associated

14    with a p53 mutation, wherein the composition

15    comprises (a) a c-FLIP inhibitor

16    (b) a chemotherapeutic agent, wherein the

17    chemotherapeutic agent is a thymidylate synthase

18    inhibitor, a platinum cytotoxic agent or a

19    topoisomerase inhibitor

20    and

21    (c) a pharmaceutically acceptable excipient, diluent

22    or carrier.

23

24    A fifth aspect provides a kit for the treatment of

25    cancer associated with a p53 mutation, said kit

26    comprising

27    (a) a c-FLIP inhibitor and

28    (b) a chemotherapeutic agent, wherein the

29    chemotherapeutic agent is a thymidylate synthase

30    inhibitor, a platinum cytotoxic agent or a

31    topoisomerase inhibitor and

32    (c) instructions for the administration of (a) and

1 (b) separately, sequentially or simultaneously.

2

3 In any of the first to fifth aspects of the  
4 invention, the c-FLIP inhibitor and the  
5 chemotherapeutic agent may be provided and  
6 administered in the absence of other active agents.  
7 However, in a preferred embodiment of these aspects  
8 aspects of the invention, there is provided (c) a  
9 death receptor binding member, or a nucleic acid  
10 encoding said binding member.

11

12 Any suitable death receptor binding member may be  
13 used. Death receptors include, Fas, TNFR, DR-3, DR-4  
14 and DR-5. In preferred embodiments of the invention,  
15 the death receptor is FAS.

16

17 The c-FLIP inhibitor, the chemotherapeutic agent  
18 and where applicable the death receptor ligand, may  
19 be administered simultaneously, sequentially or  
20 simultaneously. In preferred embodiments of the  
21 invention, the c-FLIP inhibitor is administered  
22 prior to the chemotherapeutic agent and, where  
23 applicable, the specific binding member.

24

25 A preferred binding member for use in the invention  
26 is an antibody or a fragment thereof. In  
27 particularly preferred embodiments, the binding  
28 member is the FAS antibody CH11 (Yonehara, S.,  
29 Ishii, A. and Yonehara, M. (1989) J. Exp. Med. 169,  
30 1747-1756) (available commercially e.g. from Upstate  
31 Biotechnology, Lake Placid, NY).

32

1 Any suitable thymidylate synthase inhibitor,  
2 platinum cytotoxic agent or topoisomerase inhibitor  
3 may be used in the present invention. Examples of  
4 thymidylate synthase inhibitors which may be used in  
5 the methods of the invention include 5-FU, MTA and  
6 TDX. In a preferred embodiment, the thymidylate  
7 synthase inhibitor is 5-FU. Examples of platinum  
8 cytotoxic agents which may be used include cisplatin  
9 and oxaliplatin. In a particularly preferred  
10 embodiment of the invention, the chemotherapeutic  
11 agent is cisplatin. Any suitable topoisomerase  
12 inhibitor may be used in the present invention. In  
13 a preferred embodiment, the topoisomerase inhibitor  
14 is a topoisomerase I inhibitor, for example a  
15 camptothecin. A suitable topoisomerase I inhibitor,  
16 which may be used in the present invention is  
17 irenotecan (CPT-11). Unless, the context demand  
18 otherwise, reference to CPT-11 should be taken to  
19 encompass CPT-11 or its active metabolite SN-38.  
20

21 In preferred embodiments of the invention, the c-  
22 FLIP inhibitor and the chemotherapeutic agent are  
23 administered in a potentiating ratio. the term  
24 "potentiating ratio" in the context of the present  
25 invention is used to indicate that the cFLIP  
26 inhibitor and chemotherapeutic agent are present in  
27 a ratio such that the cytotoxic activity of the  
28 combination is greater than that of either component  
29 alone or of the additive activity that would be  
30 predicted for the combinations based on the  
31 activities of the individual components. Thus in a



1     potentiating ratio, the individual components act  
2     synergistically.

3  
4     Synergism may be defined using a number of methods.  
5     For example, synergism may be defined as an RI of  
6     greater than unity using the method of Kern as  
7     modified by Romaneli (1998a, 1998b). The RI may be  
8     calculated as the ratio of expected cell survival  
9     ( $S_{exp}$ , defined as the product of the survival  
10    observed with drug A alone and the survival observed  
11    with drug B alone) to the observed cell survival  
12    ( $S_{obs}$ ) for the combination of A and B ( $RI = S_{exp}/S_{obs}$ ).  
13    Synergism may then be defined as an RI of greater  
14    than unity.

15  
16    In another method, synergism may be determined by  
17    calculating the combination index (CI) according to  
18    the method of Chou and Talalay. CI values of 1, <1,  
19    and >1 indicate additive, synergistic and  
20    antagonistic effects respectively.

21  
22    In a preferred embodiment of the invention, the c-  
23    FLIP inhibitor and the chemotherapeutic agent are  
24    present in concentrations sufficient to produce a CI  
25    of less than 1, preferably less than 0.85.

26  
27    Synergism is preferably defined as an RI of greater  
28    than unity using the method of Kern as modified by  
29    Romaneli (1998a,b)). The RI may be calculated as the  
30    ratio of expected cell survival ( $S_{exp}$ , defined as the  
31    product of the survival observed with drug A alone  
32    and the survival observed with drug B alone) to the

1 observed cell survival ( $S_{obs}$ ) for the combination of  
2 A and B ( $RI = S_{exp} / S_{obs}$ ). Synergism may then be defined  
3 as an RI of greater than unity.

4  
5 In preferred embodiments of the invention, said  
6 specific binding member and chemotherapeutic agent  
7 are provided in concentrations sufficient to produce  
8 an RI of greater than 1.5, more preferably greater  
9 than 2.0, most preferably greater than 2.25.

10  
11 The combined medicament thus preferably produces a  
12 synergistic effect when used to treat tumour cells.

13  
14 The invention according to any of the first, second  
15 third, fourth and fifth aspect of the invention may  
16 be used for the killing of any cancer cell having a  
17 p53 mutation. The mutation may partially or totally  
18 inactivate p53 in a cell. In one embodiment of the  
19 invention, the p53 mutation is a p53 mutation, which  
20 totally inactivates p53. In another embodiment, the  
21 p53 mutation is a missense mutation resulting in the  
22 substitution of histidine (R175H mutation). In  
23 another embodiment, the p53 mutation is a missense  
24 mutation resulting in the substitution of tryptophan  
25 (R248W mutation) for arginine.

26  
27 As described in the Examples, as well as testing the  
28 cytotoxicity of combinations of c-FLIP inhibitors  
29 and chemotherapeutic agents on cancer cells, the  
30 inventors further tested the effects of c-FLIP  
31 alone. The inventors unexpectedly observed that  
32 relatively potent inhibition of cFLIP using high

1 concentrations of siRNA triggered apoptosis in the  
2 absence of chemotherapy in both RKO and H630 cell  
3 lines. This demonstration that cFLIP inhibition in  
4 the absence of chemotherapy is sufficient to trigger  
5 apoptosis in cancer cells enables the use of c-FLIP  
6 inhibition alone as a chemotherapeutic strategy.

7  
8 Accordingly, in a sixth aspect of the invention,  
9 there is provided a method of killing cancer cells,  
10 comprising administration to said cells of an  
11 effective amount of a c-FLIP inhibitor, wherein the  
12 c-FLIP inhibitor is administered as the sole  
13 cytotoxic agent in the substantial absence of other  
14 cytotoxic agents.

15  
16 A seventh aspect of the invention provides a method  
17 of treating cancer comprising administration to a  
18 subject in need thereof a therapeutically effective  
19 amount of a c-FLIP inhibitor, wherein the c-FLIP  
20 inhibitor is administered as the sole cytotoxic  
21 agent in the substantial absence of other cytotoxic  
22 agents.

23  
24 An eighth aspect provides the use of a c-FLIP  
25 inhibitor as the sole cytotoxic agent in the  
26 preparation of a medicament for treating cancer,  
27 wherein the medicament is for treatment in the  
28 substantial absence of other cytotoxic agents.

29  
30 A ninth aspect provides a pharmaceutical composition  
31 for the treatment of cancer, wherein the composition  
32 comprises a c-FLIP inhibitor as the sole cytotoxic

1 agent and a pharmaceutically acceptable excipient,  
2 diluent or carrier, wherein the composition is for  
3 treatment in the absence of other cytotoxic agents.

4  
5 The sixth to ninth aspects of the invention may be  
6 used in the treatment of any cancer. The cancer  
7 cells may comprise a p53 wild type genotype or,  
8 alternatively, may comprise p53 mutant genotypes.  
9 The mutation may partially or totally inactivate p53  
10 in a cell. In one embodiment of the invention, the  
11 p53 mutation is a p53 mutation, which totally  
12 inactivates p53. In another embodiment, the p53  
13 mutation is a missense mutation resulting in the  
14 substitution of histidine (R175H mutation). In  
15 another embodiment, the p53 mutation is a missense  
16 mutation resulting in the substitution of tryptophan  
17 (R248W mutation) for arginine.

18  
19 Any suitable c-FLIP inhibitor may be used in methods  
20 of the invention. The inhibitor may be peptide or  
21 non-peptide.

22  
23 In one preferred embodiment, said c-FLIP inhibitor  
24 is an antisense molecule which modulates the  
25 expression of the gene encoding c-FLIP.

26  
27 In a more preferred embodiment, said c-FLIP  
28 inhibitor is an RNAi agent, which modulates  
29 expression of the c-FLIP gene. The agent may be an  
30 siRNA, an shRNA, a ddRNAi construct or a  
31 transcription template thereof, e.g., a DNA encoding  
32 an shRNA. In preferred embodiments the RNAi agent

1 is an siRNA which is homologous to a part of the  
2 mRNA sequence of the gene encoding c-FLIP.

3

4 Preferred RNAi agents of and for use in the  
5 invention are between 15 and 25 nucleotides in  
6 length, preferably between 19 and 22 nucleotides,  
7 most preferably 21 nucleotides in length. In  
8 particularly preferred embodiments of the invention,  
9 the RNAi agent has the nucleotide sequence shown as  
10 SEQ ID NO: 1.

11

12 AAG CAG TCT GTT CAA GGA GCA (SEQ ID NO: 1)

13

14 In another particularly preferred embodiment of the  
15 invention, the RNAi agent has the nucleotide  
16 sequence shown as SEQ ID NO: 2

17

18 AAG GAA CAG CTT GGC GCT CAA (SEQ ID NO: 2)

19

20 Indeed such RNAi agents represents a tenth and  
21 eleventh independent aspects of the present  
22 invention.

23

24 According to a further aspect of the invention,  
25 there is provided a vector comprising the RNAi agent  
26 of the tenth aspect of the invention.

27

28 In a further aspect, there is provided a kit for the  
29 treatment of cancer associated with a p53 mutation,  
30 said kit comprising

31 (a) a c-FLIP inhibitor and

32 (b) a chemotherapeutic agent, wherein the

1     chemotherapeutic agent is a thymidylate synthase  
2     inhibitor, a platinum cytotoxic agent or a  
3     topoisomerase inhibitor and  
4     (c) instructions for the administration of (a) and  
5     (b) separately, sequentially or simultaneously.

6

7     Preferred features of each aspect of the invention  
8     are as for each of the other aspects mutatis  
9     mutandis unless the context demands otherwise.

10

#### 11     **Detailed Description**

12

13     As described above, the present invention relates to  
14     methods of treatment of cancer, involving cFLIP  
15     inhibition.

16

17     The methods of the invention may involve the  
18     determination of expression of FLIP protein.

19

20     The expression of FLIP may be measured using any  
21     technique known in the art. Either mRNA or protein  
22     can be measured as a means of determining up-or down  
23     regulation of expression of a gene. Quantitative  
24     techniques are preferred. However semi-quantitative  
25     or qualitative techniques can also be used. Suitable  
26     techniques for measuring gene products include, but  
27     are not limited to, SAGE analysis, DNA microarray  
28     analysis, Northern blot,  
29     Western blot, immunocytochemical analysis, and  
30     ELISA.

31

1 RNA can be detected using any of the known  
2 techniques in the art. Preferably an amplification  
3 step is used as the amount of RNA from the sample  
4 may be very small. Suitable techniques may include  
5 real-time RT-PCR, hybridisation of copy mRNA (cRNA)  
6 to an array of nucleic acid probes and Northern  
7 Blotting.

8  
9 For example, when using mRNA detection, the method  
10 may be carried out by converting the isolated mRNA  
11 to cDNA according to standard methods; treating the  
12 converted cDNA with amplification reaction reagents  
13 (such as cDNA PCR reaction reagents) in a container  
14 along with an appropriate mixture of nucleic acid  
15 primers; reacting the contents of the container to  
16 produce amplification products; and analyzing the  
17 amplification products to detect the presence of  
18 gene expression products of one or more of the genes  
19 encoding FLIP protein. Analysis may be accomplished  
20 using Southern Blot analysis to detect the presence  
21 of the gene products in the amplification product.  
22 Southern Blot analysis is known in the art. The  
23 analysis step may be further accomplished by  
24 quantitatively detecting the presence of such gene  
25 products in the amplification products, and  
26 comparing the quantity of product detected against a  
27 panel of expected values for known presence or  
28 absence in normal and malignant tissue derived using  
29 similar primers.

30  
31 In e.g. determining gene expression in carrying out  
32 conventional molecular biological, microbiological

1 and recombinant DNA techniques techniques known in  
2 the art may be employed. Details of such  
3 techniques are described in, for example, Sambrook,  
4 Fritsch and Maniatis, "Molecular Cloning, A  
5 Laboratory Manual, Cold Spring Harbor Laboratory  
6 Press, 1989, and Ausubel et al, Short Protocols in  
7 Molecular Biology, John Wiley and Sons, 1992).

8

9 **Binding members**

10

11 In the context of the present invention, a "binding  
12 member" is a molecule which has binding specificity  
13 for another molecule, in particular a receptor,  
14 preferably a death receptor. The binding member may  
15 be a member of a pair of specific binding members.  
16 The members of a binding pair may be naturally  
17 derived or wholly or partially synthetically  
18 produced. One member of the pair of molecules may  
19 have an area on its surface, which may be a  
20 protrusion or a cavity, which specifically binds to  
21 and is therefore complementary to a particular  
22 spatial and polar organisation of the other member  
23 of the pair of molecules. Thus, the members of the  
24 pair have the property of binding specifically to  
25 each other. A binding member of the invention and  
26 for use in the invention may be any moiety, for  
27 example an antibody or ligand, which preferably can  
28 bind to a death receptor.

29

30 The binding member may bind to any death receptor.  
31 Death receptors include, Fas, TNFR, DR-3, DR-4 and



1 DR-5. In preferred embodiments of the invention, the  
2 death receptor is FAS.

3

4 In preferred embodiments, the binding member  
5 comprises at least one human constant region.

6

#### 7 **Antibodies**

8

9 An "antibody" is an immunoglobulin, whether natural  
10 or partly or wholly synthetically produced. The  
11 term also covers any polypeptide, protein or peptide  
12 having a binding domain which is, or is homologous  
13 to, an antibody binding domain. These can be  
14 derived from natural sources, or they may be partly  
15 or wholly synthetically produced. Examples of  
16 antibodies are the immunoglobulin isotypes and their  
17 isotypic subclasses and fragments which comprise an  
18 antigen binding domain such as Fab, scFv, Fv, dAb,  
19 Fd; and diabodies.

20

21 A binding member for use in certain embodiments, the  
22 invention may be an antibody such as a monoclonal or  
23 polyclonal antibody, or a fragment thereof. The  
24 constant region of the antibody may be of any class  
25 including, but not limited to, human classes IgG,  
26 IgA, IgM, IgD and IgE. The antibody may belong to  
27 any sub class e.g. IgG1, IgG2, IgG3 and IgG4. IgG1  
28 is preferred.

29

30 As antibodies can be modified in a number of ways,  
31 the term "antibody" should be construed as covering  
32 any binding member or substance having a binding

1 domain with the required specificity. Thus, this  
2 term covers antibody fragments, derivatives,  
3 functional equivalents and homologues of antibodies,  
4 including any polypeptide comprising an  
5 immunoglobulin binding domain, whether natural or  
6 wholly or partially synthetic. Chimeric molecules  
7 comprising an immunoglobulin binding domain, or  
8 equivalent, fused to another polypeptide are  
9 therefore included. Cloning and expression of  
10 chimeric antibodies are described in EP-A-0120694  
11 and EP-A-0125023.

12  
13 Examples of such fragments which can be used in the  
14 invention include the Fab fragment, the Fd fragment,  
15 the Fv fragment, the dAb fragment (Ward, E.S. et  
16 al., Nature 341:544-546 (1989)), F(ab')<sub>2</sub> fragments,  
17 single chain Fv molecules (scFv), bispecific single  
18 chain Fv dimers (PCT/US92/09965) and "diabodies",  
19 multivalent or multispecific fragments constructed  
20 by gene fusion (WO94/13804; P. Hollinger et al.,  
21 Proc. Natl. Acad. Sci. USA 90:6444-6448 (1993)).

22  
23 A fragment of an antibody or of a polypeptide for  
24 use in the present invention generally means a  
25 stretch of amino acid residues of at least 5 to 7  
26 contiguous amino acids, often at least about 7 to 9  
27 contiguous amino acids, typically at least about 9  
28 to 13 contiguous amino acids, more preferably at  
29 least about 20 to 30 or more contiguous amino acids  
30 and most preferably at least about 30 to 40 or more  
31 consecutive amino acids.

32

1 A "derivative" of such an antibody or polypeptide,  
2 or of a fragment antibody means an antibody or  
3 polypeptide modified by varying the amino acid  
4 sequence of the protein, e.g. by manipulation of the  
5 nucleic acid encoding the protein or by altering the  
6 protein itself. Such derivatives of the natural  
7 amino acid sequence may involve insertion, addition,  
8 deletion and/or substitution of one or more amino  
9 acids, preferably while providing a peptide having  
10 death receptor, e.g. FAS neutralisation and/or  
11 binding activity. Preferably such derivatives  
12 involve the insertion, addition, deletion and/or  
13 substitution of 25 or fewer amino acids, more  
14 preferably of 15 or fewer, even more preferably of  
15 10 or fewer, more preferably still of 4 or fewer and  
16 most preferably of 1 or 2 amino acids only.

17

18 In preferred embodiments, the binding member is  
19 humanised. Methods for making humanised antibodies  
20 are known in the art e.g see U.S. Patent No.  
21 5,225,539. A humanised antibody may be a modified  
22 antibody having the hypervariable region of a  
23 monoclonal antibody and the constant region of a  
24 human antibody. Thus the binding member may  
25 comprise a human constant region. The variable  
26 region other than the hypervariable region may also  
27 be derived from the variable region of a human  
28 antibody and/or may also be derived from a  
29 monoclonal antibody. In such case, the entire  
30 variable region may be derived from murine  
31 monoclonal antibody and the antibody is said to be  
32 chimerised. Methods for making chimerised

1     antibodies are known in the art (e.g see U.S. Patent  
2     Nos. 4,816,397 and 4,816,567).

3

4     It is possible to take monoclonal and other  
5     antibodies and use techniques of recombinant DNA  
6     technology to produce other antibodies or chimeric  
7     molecules which retain the specificity of the  
8     original antibody. Such techniques may involve  
9     introducing DNA encoding the immunoglobulin variable  
10    region, or the complementary determining regions  
11    (CDRs), of an antibody to the constant regions, or  
12    constant regions plus framework regions, of a  
13    different immunoglobulin. See, for instance, EP-A-  
14    184187, GB 2188638A or EP-A-239400. A hybridoma or  
15    other cell producing an antibody may be subject to  
16    genetic mutation or other changes, which may or may  
17    not alter the binding specificity of antibodies  
18    produced.

19

20    A typical antibody for use in the present invention  
21    is a humanised equivalent of CH11 or any chimerised  
22    equivalent of an antibody that can bind to the FAS  
23    receptor and any alternative antibodies directed at  
24    the FAS receptor that have been chimerised and can  
25    be use in the treatment of humans. Furthermore, the  
26    typical antibody is any antibody that can cross-  
27    react with the extracellular portion of the FAS  
28    receptor and either bind with high affinity to the  
29    FAS receptor, be internalised with the FAS receptor  
30    or trigger signalling through the FAS receptor.

31

32    **Production of Binding Members**

1  
2 Binding members, which may be used in certain  
3 aspects of the present invention may be generated  
4 wholly or partly by chemical synthesis. The binding  
5 members can be readily prepared according to well-  
6 established, standard liquid or, preferably, solid-  
7 phase peptide synthesis methods, general  
8 descriptions of which are broadly available (see,  
9 for example, in J.M. Stewart and J.D. Young, Solid  
10 Phase Peptide Synthesis, 2nd edition, Pierce  
11 Chemical Company, Rockford, Illinois (1984), in M.  
12 Bodanzsky and A. Bodanzsky, The Practice of Peptide  
13 Synthesis, Springer Verlag, New York (1984); and  
14 Applied Biosystems 430A Users Manual, ABI Inc.,  
15 Foster City, California), or they may be prepared in  
16 solution, by the liquid phase method or by any  
17 combination of solid-phase, liquid phase and  
18 solution chemistry, e.g. by first completing the  
19 respective peptide portion and then, if desired and  
20 appropriate, after removal of any protecting groups  
21 being present, by introduction of the residue X by  
22 reaction of the respective carbonic or sulfonic acid  
23 or a reactive derivative thereof.  
24  
25 Another convenient way of producing a binding member  
26 suitable for use in the present invention is to  
27 express nucleic acid encoding it, by use of nucleic  
28 acid in an expression system. Thus the present  
29 invention further provides the use of (a) nucleic  
30 acid encoding a specific binding member which binds  
31 to a cell death receptor and (b) a chemotherapeutic  
32 agent and (c) a CFLIP inhibitor in the preparation

1 of a medicament for treating cancer associated with  
2 a p53 mutation.

3  
4 Nucleic acids of and/or for use in accordance with  
5 the present invention may comprise DNA or RNA and  
6 may be wholly or partially synthetic. In a preferred  
7 aspect, nucleic acid for use in the invention codes  
8 for a binding member of the invention as defined  
9 above. The skilled person will be able to determine  
10 substitutions, deletions and/or additions to such  
11 nucleic acids which will still provide a binding  
12 member suitable for use in the present invention.

13  
14 Nucleic acid sequences encoding a binding member for  
15 use with the present invention can be readily  
16 prepared by the skilled person using the information  
17 and references contained herein and techniques known  
18 in the art (for example, see Sambrook, Fritsch and  
19 Maniatis, "Molecular Cloning", A Laboratory Manual,  
20 Cold Spring Harbor Laboratory Press, 1989, and  
21 Ausubel et al, Short Protocols in Molecular Biology,  
22 John Wiley and Sons, 1992), given the nucleic acid  
23 sequences and clones available. These techniques  
24 include (i) the use of the polymerase chain reaction  
25 (PCR) to amplify samples of such nucleic acid, e.g.  
26 from genomic sources, (ii) chemical synthesis, or  
27 (iii) preparing cDNA sequences. DNA encoding  
28 antibody fragments may be generated and used in any  
29 suitable way known to those of skill in the art,  
30 including by taking encoding DNA, identifying  
31 suitable restriction enzyme recognition sites either  
32 side of the portion to be expressed, and cutting out

1     said portion from the DNA. The portion may then be  
2     operably linked to a suitable promoter in a standard  
3     commercially available expression system. Another  
4     recombinant approach is to amplify the relevant  
5     portion of the DNA with suitable PCR primers.  
6     Modifications to the sequences can be made, e.g.  
7     using site directed mutagenesis, to lead to the  
8     expression of modified peptide or to take account of  
9     codon preferences in the host cells used to express  
10    the nucleic acid.

11  
12    The nucleic acid may be comprised as construct(s) in  
13    the form of a plasmid, vector, transcription or  
14    expression cassette which comprises at least one  
15    nucleic acid as described above. The construct may  
16    be comprised within a recombinant host cell which  
17    comprises one or more constructs as above.  
18    Expression may conveniently be achieved by culturing  
19    under appropriate conditions recombinant host cells  
20    containing the nucleic acid. Following production  
21    by expression a specific binding member may be  
22    isolated and/or purified using any suitable  
23    technique, then used as appropriate.

24  
25    Binding members-encoding nucleic acid molecules and  
26    vectors for use in accordance with the present  
27    invention may be provided isolated and/or purified,  
28    e.g. from their natural environment, in  
29    substantially pure or homogeneous form, or, in the  
30    case of nucleic acid, free or substantially free of  
31    nucleic acid or genes of origin other than the

1 sequence encoding a polypeptide with the required  
2 function.

3  
4 Systems for cloning and expression of a polypeptide  
5 in a variety of different host cells are well known.  
6 Suitable host cells include bacteria, mammalian  
7 cells, yeast and baculovirus systems. Mammalian  
8 cell lines available in the art for expression of a  
9 heterologous polypeptide include Chinese hamster  
10 ovary cells, HeLa cells, baby hamster kidney cells,  
11 NSO mouse melanoma cells and many others. A common,  
12 preferred bacterial host is *E. coli*.

13  
14 The expression of antibodies and antibody fragments  
15 in prokaryotic cells such as *E. coli* is well  
16 established in the art. For a review, see for  
17 example Plückthun, *Bio/Technology* 9:545-551 (1991).  
18 Expression in eukaryotic cells in culture is also  
19 available to those skilled in the art as an option  
20 for production of a binding member, see for recent  
21 review, for example Reff, *Curr. Opinion Biotech.*  
22 4:573-576 (1993); Trill et al., *Curr. Opinion*  
23 *Biotech.* 6:553-560 (1995).

24  
25 Suitable vectors can be chosen or constructed,  
26 containing appropriate regulatory sequences,  
27 including promoter sequences, terminator sequences,  
28 polyadenylation sequences, enhancer sequences,  
29 marker genes and other sequences as appropriate.  
30 Vectors may be plasmids, viral e.g. 'phage, or  
31 phagemid, as appropriate. For further details see,  
32 for example, Sambrook et al., *Molecular Cloning: A*



1 Laboratory Manual: 2nd Edition, Cold Spring Harbor  
2 Laboratory Press (1989). Many known techniques and  
3 protocols for manipulation of nucleic acid, for  
4 example in preparation of nucleic acid constructs,  
5 mutagenesis, sequencing, introduction of DNA into  
6 cells and gene expression, and analysis of proteins,  
7 are described in detail in Ausubel et al. eds.,  
8 Short Protocols in Molecular Biology, 2nd Edition,  
9 John Wiley & Sons (1992).

10  
11 The nucleic acid may be introduced into a host cell  
12 by any suitable means. The introduction may employ  
13 any available technique. For eukaryotic cells,  
14 suitable techniques may include calcium phosphate  
15 transfection, DEAE-Dextran, electroporation,  
16 liposome-mediated transfection and transduction  
17 using retrovirus or other virus, e.g. vaccinia or,  
18 for insect cells, baculovirus. For bacterial cells,  
19 suitable techniques may include calcium chloride  
20 transformation, electroporation and transfection  
21 using bacteriophage.

22  
23 Marker genes such as antibiotic resistance or  
24 sensitivity genes may be used in identifying clones  
25 containing nucleic acid of interest, as is well  
26 known in the art.

27  
28 The introduction may be followed by causing or  
29 allowing expression from the nucleic acid, e.g. by  
30 culturing host cells under conditions for expression  
31 of the gene.

32

1 The nucleic acid may be integrated into the genome  
2 (e.g. chromosome) of the host cell. Integration may  
3 be promoted by inclusion of sequences which promote  
4 recombination with the genome in accordance with  
5 standard techniques. The nucleic acid may be on an  
6 extra-chromosomal vector within the cell, or  
7 otherwise identifiably heterologous or foreign to  
8 the cell.

9

#### 10 **RNAi agents**

11

12 As described herein, c-FLIP inhibitors for use in  
13 the invention may be RNAi agents.

14

15 RNA interference (RNAi) or posttranscriptional gene  
16 silencing (PTGS) is a process whereby double-  
17 stranded RNA induces potent and specific gene  
18 silencing. RNAi is mediated by RNA-induced silencing  
19 complex (RISC), a sequence-specific, multicomponent  
20 nuclease that destroys messenger RNAs homologous to  
21 the silencing trigger. RISC is known to contain  
22 short RNAs (approximately 22 nucleotides) derived  
23 from the double-stranded RNA trigger.

24

25 In one aspect, the invention provides methods of  
26 employing an RNAi agent to modulate expression,  
27 preferably reducing expression of a target gene, c-  
28 FLIP, in a mammalian, preferably human host. By  
29 reducing expression is meant that the level of  
30 expression of a target gene or coding sequence is  
31 reduced or inhibited by at least about 2-fold,  
32 usually by at least about 5-fold, e.g., 10-fold, 15-

1 fold, 20-fold, 50-fold, 100-fold or more, as  
2 compared to a control. In certain embodiments, the  
3 expression of the target gene is reduced to such an  
4 extent that expression of the c-FLIP gene /coding  
5 sequence is effectively inhibited. By modulating  
6 expression of a target gene is meant altering, e.g.,  
7 reducing, translation of a coding sequence, e.g.,  
8 genomic DNA, mRNA etc., into a polypeptide, e.g.,  
9 protein, product.

10

11 The RNAi agents that may be employed in preferred  
12 embodiments of the invention are small ribonucleic  
13 acid molecules (also referred to herein as  
14 interfering ribonucleic acids), that are present in  
15 duplex structures, e.g., two distinct  
16 oligoribonucleotides hybridized to each other or a  
17 single ribooligonucleotide that assumes a small  
18 hairpin formation to produce a duplex structure.  
19 Preferred oligoribonucleotides are ribonucleic  
20 acids of not greater than 100 nt in length,  
21 typically not greater than 75 nt in length. Where  
22 the RNA agent is an siRNA, the length of the duplex  
23 structure typically ranges from about 15 to 30 bp,  
24 usually from about 20 and 29 bps, most preferably 21  
25 bp. Where the RNA agent is a duplex structure of a  
26 single ribonucleic acid that is present in a hairpin  
27 formation, i.e., a shRNA, the length of the  
28 hybridized portion of the hairpin is typically the  
29 same as that provided above for the siRNA type of  
30 agent or longer by 4-8 nucleotides.

31

1 In certain embodiments, instead of the RNAi agent  
2 being an interfering ribonucleic acid, e.g., an  
3 siRNA or shRNA as described above, the RNAi agent  
4 may encode an interfering ribonucleic acid. In these  
5 embodiments, the RNAi agent is typically a DNA that  
6 encodes the interfering ribonucleic acid. The DNA  
7 may be present in a vector.

8  
9 The RNAi agent can be administered to the host using  
10 any suitable protocol known in the art. For example,  
11 the nucleic acids may be introduced into tissues or  
12 host cells by viral infection, microinjection,  
13 fusion of vesicles, particle bombardment, or  
14 hydrodynamic nucleic acid administration.

15  
16 DNA directed RNA interference (ddRNAi) is an RNAi  
17 technique which may be used in the methods of the  
18 invention. ddRNAi is described in U.S. 6,573,099 and  
19 GB 2353282. ddRNAi is a method to trigger RNAi  
20 which involves the introduction of a DNA construct  
21 into a cell to trigger the production of double  
22 stranded (dsRNA), which is then cleaved into small  
23 interfering RNA (siRNA) as part of the RNAi process.  
24 ddRNAi expression vectors generally employ RNA  
25 polymerase III promoters (e.g. U6 or H1) for the  
26 expression of siRNA target sequences transfected in  
27 mammalian cells. siRNA target sequences generated  
28 from a ddRNAi expression cassette system can be  
29 directly cloned into a vector that does not contain  
30 a U6 promoter. Alternatively short single stranded  
31 DNA oligos containing the hairpin siRNA target  
32 sequence can be annealed and cloned into a vector

1 downstream of the pol III promoter. The primary  
2 advantages of ddRNAi expression vectors is that they  
3 allow for long term interference effects and  
4 minimise the natural interferon response in cells..  
5

6 **Antisense nucleic acids**  
7

8 As described herein, c-FLIP inhibitors for use in  
9 the invention may be anti-sense molecules or nucleic  
10 acid constructs that express such anti-sense  
11 molecules as RNA. The antisense molecules may be  
12 natural or synthetic. Synthetic antisense molecules  
13 may have chemical modifications from native nucleic  
14 acids. The antisense sequence is complementary to  
15 the mRNA of the targeted c-FLIP gene, and inhibits  
16 expression of the targeted gene products. Antisense  
17 molecules inhibit gene expression through various  
18 mechanisms, e.g. by reducing the amount of mRNA  
19 available for translation, through activation of  
20 RNase H, or steric hindrance. One or a combination  
21 of antisense molecules may be administered, where a  
22 combination may comprise multiple different  
23 sequences.  
24

25 Antisense molecules may be produced by expression of  
26 all or a part of the c-FLIP sequence in an  
27 appropriate vector, where the transcriptional  
28 initiation is oriented such that an antisense strand  
29 is produced as an RNA molecule. Alternatively, the  
30 antisense molecule may be a synthetic  
31 oligonucleotide. Antisense oligonucleotides will  
32 generally be at least about 7, usually at least

1     about 12, more usually at least about 16 nucleotides  
2     in length, and usually not more than about 50,  
3     preferably not more than about 35 nucleotides in  
4     length.

5  
6     A specific region or regions of the endogenous c-  
7     FLIP sense strand mRNA sequence is chosen to be  
8     complemented by the antisense sequence. Selection of  
9     a specific sequence for the oligonucleotide may use  
10    an empirical method, where several candidate  
11    sequences are assayed for inhibition of expression  
12    of the target gene in an in vitro or animal model. A  
13    combination of sequences may also be used, where  
14    several regions of the mRNA sequence are selected  
15    for antisense complementation.

16  
17    Antisense oligonucleotides may be chemically  
18    synthesized by methods known in the art (see Wagner  
19    et al. (1993), supra, and Milligan et al., supra.)  
20    Preferred oligonucleotides are chemically modified  
21    from the native phosphodiester structure, in order  
22    to increase their intracellular stability and  
23    binding affinity. A number of such modifications  
24    have been described in the literature, which alter  
25    the chemistry of the backbone, sugars or  
26    heterocyclic bases. Among useful changes in the  
27    backbone chemistry are phosphorodiamidate linkages,  
28    methylphosphonates phosphorothioates;  
29    phosphorodithioates, where both of the non-bridging  
30    oxygens are substituted with sulfur;  
31    phosphoroamidites; alkyl phosphotriesters and  
32    boranophosphates. Achiral phosphate derivatives

1 include 3'-O-5'-S-phosphorothioate, 3'-S-5'-O-  
2 phosphorothioate, 3'-CH<sub>2</sub>-5'-O-phosphonate and 3'-NH-  
3 5'-O-phosphoroamidate. Peptide nucleic acids may  
4 replace the entire ribose phosphodiester backbone  
5 with a peptide linkage. Sugar modifications may also  
6 be used to enhance stability and affinity.

7

### 8 **Chemotherapeutic Agents**

9 Any suitable thymidylate synthase inhibitor,  
10 platinum cytotoxic agent or topoisomerase inhibitor  
11 may be used in the present invention. Examples of  
12 thymidylate synthase inhibitors which may be used in  
13 the methods of the invention include 5-FU, MTA and  
14 TDX. In a preferred embodiment, the thymidylate  
15 synthase inhibitor is 5-FU. Examples of platinum  
16 cytotoxic agents which may be used include cisplatin  
17 and oxaliplatin. In a particularly preferred  
18 embodiment of the invention, the chemotherapeutic  
19 agent is cisplatin. A topoisomerase inhibitor, which  
20 may be used in the present invention is irenotecan  
21 (CPT-11).

22

### 23 **Treatment**

24

25 "Treatment" includes any regime that can benefit a  
26 human or non-human animal. The treatment may be in  
27 respect of an existing condition or may be  
28 prophylactic (preventative treatment). Treatment may  
29 include curative, alleviation or prophylactic  
30 effects.

31

1 "Treatment of cancer" includes treatment of  
2 conditions caused by cancerous growth and includes  
3 the treatment of neoplastic growths or tumours.  
4 Examples of tumours that can be treated using the  
5 invention are, for instance, sarcomas, including  
6 osteogenic and soft tissue sarcomas, carcinomas,  
7 e.g., breast-, lung-, bladder-, thyroid-, prostate-,  
8 colon-, rectum-, pancreas-, stomach-, liver-,  
9 uterine-, cervical and ovarian carcinoma, lymphomas,  
10 including Hodgkin and non-Hodgkin lymphomas,  
11 neuroblastoma, melanoma, myeloma, Wilms tumor, and  
12 leukemias, including acute lymphoblastic leukaemia  
13 and acute myeloblastic leukaemia, gliomas and  
14 retinoblastomas.

15

16 In preferred embodiments of the invention, the  
17 cancer is one or more of colorectal, breast ,  
18 ovarian, cervical, gastric, lung, liver, skin and  
19 myeloid (e.g. bone marrow) cancer.

20

## 21 **Administration**

22

23 As described above, c-FLIP inhibitors of and for use  
24 in the present invention may be administered in any  
25 suitable way. Moreover in any of the first to fifth  
26 aspects of the invention, they may be used in  
27 combination therapy with other treatments, for  
28 example, other chemotherapeutic agents or binding  
29 members. In such embodiments, the c-FLIP inhibitors  
30 or compositions of the invention may be administered  
31 simultaneously, separately or sequentially with  
32 another chemotherapeutic agent.



1

2 Where administered separately or sequentially, they  
3 may be administered within any suitable time period  
4 e.g. within 1, 2, 3, 6, 12, 24, 48 or 72 hours of  
5 each other. In preferred embodiments, they are  
6 administered within 6, preferably within 2, more  
7 preferably within 1, most preferably within 20  
8 minutes of each other.

9

10 In a preferred embodiment, the c-FLIP inhibitors  
11 and/or compositions of the invention are  
12 administered as a pharmaceutical composition, which  
13 will generally comprise a suitable pharmaceutical  
14 excipient, diluent or carrier selected dependent on  
15 the intended route of administration.

16

17 The c-FLIP inhibitors and/or compositions of the  
18 invention may be administered to a patient in need  
19 of treatment via any suitable route.

20

21 Some suitable routes of administration include (but  
22 are not limited to) oral, rectal, nasal, topical  
23 (including buccal and sublingual), vaginal or  
24 parenteral (including subcutaneous, intramuscular,  
25 intravenous, intradermal, intrathecal and epidural)  
26 administration. Intravenous administration is  
27 preferred.

28

29 The C-FLIP inhibitor, product or composition may be  
30 administered in a localised manner to a tumour site  
31 or other desired site or may be delivered in a  
32 manner in which it targets tumour or other cells.

1 Targeting therapies may be used to deliver the  
2 active agents more specifically to certain types of  
3 cell, by the use of targeting systems such as  
4 antibody or cell specific ligands. Targeting may be  
5 desirable for a variety of reasons, for example if  
6 the agent is unacceptably toxic, or if it would  
7 otherwise require too high a dosage, or if it would  
8 not otherwise be able to enter the target cells.

9  
10 For intravenous, injection, or injection at the site  
11 of affliction, the active ingredient will be in the  
12 form of a parenterally acceptable aqueous solution  
13 which is pyrogen-free and has suitable pH,  
14 isotonicity and stability. Those of relevant skill  
15 in the art are well able to prepare suitable  
16 solutions using, for example, isotonic vehicles such  
17 as Sodium Chloride Injection, Ringer's Injection,  
18 Lactated Ringer's Injection. Preservatives,  
19 stabilisers, buffers, antioxidants and/or other  
20 additives may be included, as required.

21  
22 Pharmaceutical compositions for oral administration  
23 may be in tablet, capsule, powder or liquid form. A  
24 tablet may comprise a solid carrier such as gelatin  
25 or an adjuvant. Liquid pharmaceutical compositions  
26 generally comprise a liquid carrier such as water,  
27 petroleum, animal or vegetable oils, mineral oil or  
28 synthetic oil. Physiological saline solution,  
29 dextrose or other saccharide solution or glycols  
30 such as ethylene glycol, propylene glycol or  
31 polyethylene glycol may be included.

32

1 The c-FLIP inhibitors and/or compositions of the  
2 invention may also be administered via microspheres,  
3 liposomes, other microparticulate delivery systems  
4 or sustained release formulations placed in certain  
5 tissues including blood. Suitable examples of  
6 sustained release carriers include semipermeable  
7 polymer matrices in the form of shared articles,  
8 e.g. suppositories or microcapsules. Implantable or  
9 microcapsular sustained release matrices include  
10 polylactides (US Patent No. 3, 773, 919; EP-A-  
11 0058481) copolymers of L-glutamic acid and gamma  
12 ethyl-L-glutamate (Sidman et al, Biopolymers 22(1):  
13 547-556, 1985), poly (2-hydroxyethyl-methacrylate)  
14 or ethylene vinyl acetate (Langer et al, J. Biomed.  
15 Mater. Res. 15: 167-277, 1981, and Langer, Chem.  
16 Tech. 12:98-105, 1982). Liposomes containing the  
17 polypeptides are prepared by well-known methods: DE  
18 3,218, 121A; Epstein et al, PNAS USA, 82: 3688-3692,  
19 1985; Hwang et al, PNAS USA, 77: 4030-4034, 1980;  
20 EP-A-0052522; E-A-0036676; EP-A-0088046; EP-A-  
21 0143949; EP-A-0142541; JP-A-83-11808; US Patent Nos  
22 4,485,045 and 4,544,545. Ordinarily, the liposomes  
23 are of the small (about 200-800 Angstroms)  
24 unilamellar type in which the lipid content is  
25 greater than about 30 mol. % cholesterol, the  
26 selected proportion being adjusted for the optimal  
27 rate of the polypeptide leakage.

28

29 Examples of the techniques and protocols mentioned  
30 above and other techniques and protocols which may  
31 be used in accordance with the invention can be

1 found in Remington's Pharmaceutical Sciences, 16th  
2 edition, Oslo, A. (ed), 1980.

3

4

#### 5 **Pharmaceutical Compositions**

6

7 Pharmaceutical compositions according to the present  
8 invention, and for use in accordance with the  
9 present invention may comprise, in addition to  
10 active ingredients, a pharmaceutically acceptable  
11 excipient, carrier, buffer stabiliser or other  
12 materials well known to those skilled in the art.  
13 Such materials should be non-toxic and should not  
14 interfere with the efficacy of the active  
15 ingredient. The precise nature of the carrier or  
16 other material will depend on the route of  
17 administration, which may be oral, or by injection,  
18 e.g. intravenous.

19

20 The formulation may be a liquid, for example, a  
21 physiologic salt solution containing non-phosphate  
22 buffer at pH 6.8-7.6, or a lyophilised powder.

23

#### 24 **Dose**

25

26 The c-FLIP inhibitors or compositions of the  
27 invention are preferably administered to an  
28 individual in a "therapeutically effective amount",  
29 this being sufficient to show benefit to the  
30 individual. The actual amount administered, and  
31 rate and time-course of administration, will depend  
32 on the nature and severity of what is being treated.

1 Prescription of treatment, e.g. decisions on dosage  
2 etc, is ultimately within the responsibility and at  
3 the discretion of general practitioners and other  
4 medical doctors, and typically takes account of the  
5 disorder to be treated, the condition of the  
6 individual patient, the site of delivery, the method  
7 of administration and other factors known to  
8 practitioners.

9

10

#### 11 **Brief Description of the Figures**

12

13 The invention will now be described further in the  
14 following non-limiting examples. Reference is made  
15 to the accompanying drawings in which:

16

17 Figure 1A illustrates Western blot analysis of Fas,  
18 FasL, procaspase 8, FADD, BID, Bcl-2, c-FLIP<sub>L</sub>, c-  
19 FLIP<sub>S</sub>, DcR3 and  $\beta$ -tubulin in MCF-7 cells 72 hours  
20 after treatment with 5 $\mu$ M 5-FU and 50nM TDX.

21

22 Figure 1B illustrates analysis of the interaction  
23 between Fas and FasL following treatment with 5 $\mu$ M 5-  
24 FU and 50nM TDX for 48 hours. Lysates were  
25 immunoprecipitated using a FasL polyclonal antibody  
26 and analysed by Western blot using a Fas monoclonal  
27 antibody.

28

29 Figure 1C illustrates analysis of the interaction  
30 between Fas and p43- c-FLIP<sub>L</sub> following treatment  
31 with 5 $\mu$ M 5-FU and 50nM TDX for 48 hours. Lysates  
32 were immunoprecipitated using the anti-Fas CH-11

1 monoclonal antibody and analysed by Western blot  
2 using a c-FLIP monoclonal antibody.

3  
4 Figure 2A illustrates flow cytometry of MCF-7 cells  
5 treated with no drug (control), CH-11 alone  
6 (250ng/ml), 5-FU alone (5 $\mu$ M) for 96 hours, or co-  
7 treated with 5-FU for 72 hours followed by CH-11 for  
8 a further 24 hours.

9  
10 Figure 2B illustrates flow cytometry of MCF-7 cells  
11 treated with no drug (control), CH-11 alone  
12 (250ng/ml), TDX alone (50nM) for 96 hours, or co-  
13 treated with TDX for 72 hours followed by CH-11 for  
14 a further 24 hours.

15  
16 Figure 2C illustrates Western blot analysis of Fas  
17 expression in MCF-7 cells treated with 5 $\mu$ M 5-FU for  
18 48 hours.  $\beta$ -tubulin was assessed as a loading  
19 control.

20  
21 Figure 2D illustrates flow cytometry of MCF-7 cells  
22 treated with no drug (control), CH-11 alone  
23 (250ng/ml), OXA alone (5 $\mu$ M) for 96 hours, or co-  
24 treated with OXA for 72 hours followed by CH-11 for  
25 a further 24 hours.

26  
27 Figure 2E illustrates Western blot analysis of Fas,  
28 procaspase 8 and PARP expression in MCF-7 cells  
29 treated with 5 $\mu$ M 5-FU alone for 96 hours, or co-  
30 treated with 5-FU for 72 hours followed by CH-11 for  
31 a further 24 hours.

32

1 Figure 2F illustrates Western blot analysis  
2 examining the kinetics of caspase 8 activation and  
3 c-FLIP<sub>L</sub> processing in MCF-7 cells treated for 72  
4 hours with 5 $\mu$ M 5-FU followed by 250ng/ml CH-11 for  
5 the indicated times.

6  
7 Figure 3A illustrates Western blot analysis of Fas  
8 expression in HCT116 cells treated with 5-FU, TDX or  
9 OXA for 48 hours. Equal loading was assessed using a  
10  $\beta$ -tubulin antibody.

11  
12 Figure 3B illustrates Western blot analysis of  
13 procaspase 8 and PARP expression in HCT116 cells  
14 treated no drug (Con), 5 $\mu$ M 5-FU, 100nM TDX or 2 $\mu$ M  
15 OXA in the presence or absence of co-treatment with  
16 200ng/ml CH-11. For each combined treatment the  
17 cells were pre-treated with chemotherapeutic drug  
18 for 24 hours followed by CH-11 for a further 24  
19 hours.

20  
21 Figure 4A illustrates Western blot of c-FLIP<sub>L</sub>  
22 expression in MCF-7 cells stably transfected with a  
23 FLIPL (FL) construct or empty vector (EV).

24  
25 Figure 4B illustrates MTT cell viability assays in  
26 EV68, FL44 and FL64 cells treated with 5 $\mu$ M 5-FU in  
27 combination with 250ng/ml CH-11. The combined  
28 treatment resulted in a synergistic decrease in cell  
29 viability in EV68 cells (RI=2.06), but not FL44  
30 (RI=1.14) or FL64 (1.01) cells.

31

1 Figure 4C illustrates Western blot analysis of c-  
2 FLIP<sub>L</sub>, procaspase 8 and PARP expression in EV68 and  
3 FL64 cells treated with no drug (Con) or 5 $\mu$ M 5-FU in  
4 the presence (+) or absence (-) of co-treatment with  
5 250ng/ml CH-11. For each combined treatment, the  
6 cells were pre-treated with 5-FU for 72 hours  
7 followed by CH-11 for a further 24 hours.

8  
9 Figure 5A illustrates MTT cell viability assays in  
10 EV68, FL44 and FL64 cells treated with 50nM TDX or  
11 500nM MTA in the presence and absence of 250ng/ml  
12 CH-11. Combined TDX/CH-11 treatment resulted in a  
13 synergistic decrease in cell viability in EV68 cells  
14 (RI=1.75), that was significantly reduced in FL44  
15 (RI=1.22) or FL64 (RI=1.19) cells. Combined MTA/CH-  
16 11 treatment resulted in a synergistic decrease in  
17 cell viability in EV68 cells (RI=1.86), that was  
18 significantly reduced in FL44 (RI=1.29) and FL64  
19 (RI=1.06) cells.

20  
21 Figure 5B illustrates MTT cell viability assays in  
22 EV68, FL44 and FL64 cells treated with 2.5 $\mu$ M OXA in  
23 the presence and absence of 250ng/ml CH-11. Combined  
24 OXA/CH-11 treatment resulted in a synergistic  
25 decrease in cell viability in EV68 cells (RI=2.13),  
26 that was significantly reduced in FL64 (RI=1.22) or  
27 FL44 (1.19) cells.

28  
29 Figure 5C Western blot analysis of procaspase 8 and  
30 PARP expression in EV68 and FL64 cells treated with  
31 50nM TDX or 500nM MTA in the presence (+) or absence  
32 (-) of co-treatment with 250ng/ml CH-11.



1

2 Figure 5D illustrates Western blot analysis of  
3 procaspase 8 and PARP expression in EV68 and FL64  
4 cells treated with 2.5 $\mu$ M OXA in the presence (+) or  
5 absence (-) of co-treatment with 250ng/ml CH-11. For  
6 each combined treatment, the cells were pre-treated  
7 with 5-FU for 72 hours followed by CH-11 for a  
8 further 24 hours.

9

10 Figure 6A illustrates c-FLIP<sub>L</sub> and c-FLIP<sub>S</sub> expression  
11 in HCT116 cells transfected with 0, 1 and 10nM FLIP-  
12 targeted siRNA for 48 hours. Equal loading was  
13 assessed using a  $\beta$ -tubulin antibody.

14

15 Figure 6B illustrates MTT cell viability assays of  
16 HCT116 cells transfected with 5nM FLIP-targeted (FT)  
17 or scrambled control (SC) siRNA in the presence and  
18 absence of co-treatment with 5 $\mu$ M 5-FU. Combined  
19 treatment with 5-FU and FT siRNA resulted in a  
20 synergistic decrease in cell viability (RI=1.92,  
21  $p<0.0005$ ). No synergistic decrease in viability was  
22 observed in cells co-treated with 5-FU and SC siRNA  
23 (RI=0.98).

24

25 Figure 6C illustrates Western blot analysis of  
26 caspase 8 activation and PARP cleavage in HCT116  
27 cells 48 hours after treatment with no drug, 5 $\mu$ M 5-  
28 FU or 100nM TDX in mock transfected cells (M), cells  
29 transfected with 1nM scrambled control (SC) and  
30 cells transfected with 1nM FLIP-targeted (FT) siRNA.

31

1 Figure 7A illustrates c-FLIP<sub>L</sub> and c-FLIP<sub>S</sub> expression  
2 in MCF-7 cells transfected with 10nM FLIP-targeted  
3 (FT) or scrambled control (SC) siRNA for 48 hours.  
4 Equal loading was assessed using a  $\beta$ -tubulin  
5 antibody.

6  
7 Figure 7B illustrates MTT cell viability assays of  
8 MCF-7 cells transfected with 2.5nM FT siRNA in the  
9 presence and absence of co-treatment with 5 $\mu$ M 5-FU.  
10 The combined treatment resulted in a synergistic  
11 decrease in cell viability (RI=1.56, p<0.005).  
12 Figure 7C Western blot analysis of PARP cleavage in  
13 MCF-7 cells 96 hours after treatment with 5-FU in  
14 the presence (+) and absence (-) of 10nM FLIP-  
15 targeted siRNA.

16  
17 Figure 8 illustrates MTT cell viability assays of  
18 HCT116 cells transfected with 0.5nM FT or SC siRNA  
19 in the presence and absence of co-treatment with:  
20 Fig 8A 5 $\mu$ M 5-FU; Fig 8B 100nM TDX and Fig 8C 1 $\mu$ M  
21 OXA. Cells were assayed after 72 hours. Combined  
22 treatment with FT siRNA (but not SC siRNA) and each  
23 cytotoxic drug resulted in synergistic decreases in  
24 cell viability as indicated by the RI values  
25 (p<0.0005 for each combination).

26  
27 Figure 9 illustrates: A Western blot analysis of Fas  
28 expression in p53 wild type HCT116 cells treated  
29 with 5-FU or oxaliplatin (OXA) for 48 hours. B  
30 Western blot analysis of caspase 8 activation, PARP  
31 cleavage and c-FLIP expression in p53 wild type  
32 HCT116 cells treated with no drug (Con), 5 $\mu$ M 5-FU,

1 or 1 $\mu$ M OXA in the presence or absence of co-  
2 treatment with 200ng/mL CH-11. For each combined  
3 treatment the cells were pre-treated with  
4 chemotherapeutic drug for 24 hours followed by CH-11  
5 for a further 24 hours.

6  
7 Figure 10 illustrates: A c-FLIP<sub>L</sub> and c-FLIP<sub>S</sub>  
8 expression in HLacZ, HFL17, HFL24, HFS19 and HFS44  
9 cell lines. B Flow cytometric analysis of cell cycle  
10 arrest and apoptosis in HLacZ, HFL17, HFL24, HFS19  
11 and HFS44 cell lines 72 hours after treatment with  
12 5 $\mu$ M 5-FU, 1 $\mu$ M oxaliplatin (OXA) and 5 $\mu$ M CPT-11. C  
13 Flow cytometric analysis of HLacZ, HFL17, HFL24,  
14 HFS19 and HFS44 cells after co-treatment with  
15 50ng/mL CH-11 and 2.5 $\mu$ M 5-FU, 0.5 $\mu$ M oxaliplatin  
16 (OXA) and 1 $\mu$ M CPT-11. For each combined treatment  
17 the cells were pre-treated with chemotherapeutic  
18 drug for 24 hours followed by CH-11 for a further 24  
19 hours.

20  
21 Figure 11 illustrates: A c-FLIP<sub>L</sub> and c-FLIP<sub>S</sub>  
22 expression in p53 wild type HCT116 cells transfected  
23 with 1nM control siRNA (SC) and 1nM FLIP-targeted  
24 (FT) siRNA for 24 hours. B Flow cytometric analysis  
25 of apoptosis in HCT116 cells transfected with 0.5nM  
26 FT or 0.5nM SC siRNA. Transfected cells were co-  
27 treated with no drug, 5 $\mu$ M 5-FU, or 1 $\mu$ M oxaliplatin  
28 (OXA) for 48 hours. C (Panel 1) Western blot  
29 analysis of caspase 8 activation and PARP cleavage  
30 in HCT116 cells 48 hours after treatment of mock  
31 transfected cells (M), cells transfected with 0.5nM  
32 SC and cells transfected with 0.5nM FT siRNA with no

1 drug, 5 $\mu$ M 5-FU or 100nM TDX. (Panel 2) Caspase 8  
2 activation and PARP cleavage in HCT116 cells  
3 transfected with 0.5nM SC or 0.5nM FT siRNA and  
4 treated with no drug, or 1 $\mu$ M oxaliplatin (OXA) for  
5 24 hours. (Panel 3) Caspase 8 activation and PARP  
6 cleavage in HCT116 cells after transfection with  
7 0.5nM SC or 0.5nM FT siRNA and treatment with no  
8 drug, 2.5 $\mu$ M or 5 $\mu$ M CPT-11 for 24 hours. **D** MTT cell  
9 viability assays in HCT116p53<sup>+/+</sup> cells transfected  
10 with FT siRNA and co-treated with 5-FU, oxaliplatin  
11 (OXA) and CPT-11. Cell viability was assayed after  
12 72 hours. The nature of the interaction between the  
13 chemotherapeutic drugs and FT siRNA was determined  
14 by calculating the combination index (CI) according  
15 to the method of Chou and Talalay. CI values of 1,  
16 <1, and >1 indicate additive, synergistic and  
17 antagonistic effects respectively. Results are  
18 representative of at least 3 separate experiments.

19  
20 Figure 12 illustrates: **A** Western blot analysis of c-  
21 FLIP<sub>L</sub> and c-FLIP<sub>S</sub> expression in p53 wild type (wt)  
22 and null HCT116 cells. **B** Western blot analysis of c-  
23 FLIP<sub>L</sub> and c-FLIP<sub>S</sub> expression in HCT116p53<sup>-/-</sup> cells  
24 transfected with 1nM control siRNA (SC) and 1nM  
25 FLIP-targeted (FT) siRNA for 24 hours. **C** Flow  
26 cytometric ananlysis of apoptosis in HCT116p53<sup>-/-</sup>  
27 cells transfected with 1nM FT or 1nM SC siRNA.  
28 Transfected cells were co-treated with no drug, 5 $\mu$ M  
29 5-FU, 5 $\mu$ M oxaliplatin (OXA) or 1 $\mu$ M CPT-11 for 72  
30 hours. **D** MTT cell viability assays in HCT116p53<sup>-/-</sup>  
31 cells transfected with FT siRNA and co-treated with  
32 5-FU, oxaliplatin (OXA), and CPT-11. Cell viability

1 was assayed after 72 hours. The nature of the  
2 interaction between the chemotherapeutic drugs and  
3 FLIP-targeted siRNAs was determined by calculating  
4 the combination index (CI) according to the method  
5 of Chou and Talalay. Results are representative of  
6 at least 3 separate experiments.

7  
8 Figure 13 illustrates: **A** c-FLIP<sub>L</sub> and c-FLIP<sub>S</sub>  
9 expression in RKO and H630 cells transfected with  
10 1nM control siRNA (SC) and 1nM FLIP-targeted (FT)  
11 siRNA for 24 hours. **B** Flow cytometric analysis of  
12 apoptosis in RKO cells transfected with 2.5nM FT or  
13 2.5nM SC siRNA and H630 cells transfected with 1nM  
14 FT or 1nM SC siRNA. SiRNA-transfected RKO cells were  
15 co-treated with no drug, 5µM 5-FU, 1µM oxaliplatin  
16 (OXA) or 2.5µM CPT-11 for 72 hours. SiRNA-  
17 transfected H630 cells were co-treated with no drug,  
18 5µM 5-FU, 2.5µM oxaliplatin (OXA) or 1µM CPT-11 for  
19 72 hours. **C** MTT cell viability assays in RKO and  
20 H630 cells transfected with FT siRNA and co-treated  
21 with 5-FU, oxaliplatin (OXA), and CPT-11. Cell  
22 viability was assayed after 72 hours. The nature of  
23 the interaction between the chemotherapeutic drugs  
24 and FLIP-targeted siRNAs was determined by  
25 calculating the combination index (CI) according to  
26 the method of Chou and Talalay. Results are  
27 representative of at least 3 separate experiments.

28  
29 Figure 14 illustrates: **A** MTT cell viability assays  
30 in HCT116p53<sup>+/+</sup> cells transfected with FT or SC siRNA  
31 for 72 hours. **B** Western blot analysis of c-FLIP  
32 expression and PARP cleavage in p53 wild type

1 (p53<sup>+/+</sup>) and p53 null (p53<sup>-/-</sup>) HCT116 cells 24 hours  
2 after transfection with 0, 1 and 10nM FT siRNA. **C**  
3 Flow cytometric analysis of apoptosis in p53 wild  
4 type (p53<sup>+/+</sup>) and p53 null (p53<sup>-/-</sup>) HCT116 cells  
5 transfected with FT or SC siRNA for 48 hours. **D**  
6 Apoptosis in HCT116p53<sup>-/-</sup> cells transfected with FT  
7 siRNA for 48 and 72 hours. **E** Apoptosis in RKO cells  
8 transfected with FT or SC siRNA for 72 hours. **F**  
9 Apoptosis in H630 cells transfected with FT or SC  
10 siRNA for 72 hours.

11

12 Figure 15 illustrates: **A** Kinetics of c-FLIP down-  
13 regulation, caspase 8 activation and PARP cleavage  
14 in HCT116p53<sup>+/+</sup> cells transfected with 0, 1 and 10nM  
15 FT siRNA. **B** Flow cytometric analysis of the kinetics  
16 of apoptosis induction in HCT116p53<sup>+/+</sup> cells  
17 transfected with 10nM FT or 10nM SC siRNA.

18

19 Figure 16 illustrates: **A** c-FLIP<sub>L</sub> and c-FLIP<sub>S</sub>  
20 expression and PARP cleavage in p53 wild type HCT116  
21 cells transfected with 10nM control siRNA (SC) and  
22 10nM FLIP<sub>L</sub>-specific (FL) siRNA for 24 hours. **B**  
23 Western blot analysis of PARP cleavage in HCT116  
24 cells transfected with 0.5nM SC or 0.5nM FL siRNA  
25 and treated with no drug, 1μM oxaliplatin (OXA) or  
26 2.5μM for 24 hours, or 5μM 5-FU for 48 hours. **C** MTT  
27 cell viability assays in HCT116p53<sup>+/+</sup> cells  
28 transfected with FL siRNA and co-treated with 5-FU  
29 oxaliplatin (OXA), and CPT-11. Cell viability was  
30 assayed after 72 hours. The nature of the  
31 interaction between the chemotherapeutic drugs and  
32 FLIP-targeted siRNAs was determined by calculating

1 the combination index (CI) according to the method  
2 of Chou and Talalay. Results are representative of  
3 at least 3 separate experiments.

4  
5 Figure 17 illustrates illustrates graphs of RI  
6 values calculated from MTT cell viability assays of  
7 the chemotherapeutic agents 5-FU, Tomudex (TDX),  
8 CPT-11 and Oxaliplatin used in combination with the  
9 agonistic anti-Fas antibody CH-11 (200ng/ml). The RI  
10 is calculated as ratio of the expected cell survival  
11 ( $S_{exp}$ , defined as the product of the survival  
12 observed with drug A alone and the survival observed  
13 with drug B alone) to the observed cell survival  
14 ( $S_{obs}$ ) for the combination of A and B  
15 ( $RI = S_{exp}/S_{obs}$ ). Synergism is defined as an RI  
16 greater than 1.

17  
18 Figure 18 illustrates A, Flow cytometry analysis of  
19 cells stained with propidium iodide stained HCT116  
20 p53 wild-type and null cells treated with 5-FU  
21 (5 $\mu$ M), TDX (50nM), CPT-11 (5 $\mu$ M) and Oxaliplatin (1 $\mu$ M)  
22 for 24 hours and then with CH-11 (50ng/ml) for an  
23 additional 24 hours. B, Sub G0/G1 populations for  
24 the HCT116p53 wild-type and null cell lines treated  
25 with chemotherapy drugs with and without CH-11 50  
26 ng/ml.

27  
28 Figure 19 illustrates the effect of adding CH-11  
29 200ng/ml for 24 hours to HCT116 p53 wild-type and  
30 null cells already treated for 24 hours with 5-FU  
31 (5 $\mu$ M), CPT-11 (5 $\mu$ M) and Oxaliplatin (1 $\mu$ M) on PARP

1 cleavage and activation of procaspase 8 by Western  
2 blot analysis.

3

4

5 **Examples**

6

7 **MATERIALS AND METHODS**

8 **Cell Culture.** All cells were maintained in 5% CO<sub>2</sub> at  
9 37°C. MCF-7 cells were maintained in DMEM with 10%  
10 dialyzed bovine calf serum supplemented with 1mM  
11 sodium pyruvate, 2mM L-glutamine and 50µg/ml  
12 penicillin/streptomycin (from Life Technologies  
13 Inc., Paisley, Scotland). HCT116p53<sup>+/+</sup> and HCT116p53<sup>-/-</sup>  
14 isogenic human colorectal cancer cells were kindly  
15 provided by Professor Bert Vogelstein (John Hopkins  
16 University, Baltimore, MD). HCT116 cells were grown  
17 in McCoy's 5A medium (GIBCO) supplemented with 10%  
18 dialysed foetal calf serum, 50mg/ml penicillin-  
19 streptomycin, 2mM L-glutamine and 1mM sodium  
20 pyruvate. Stably transfected MCF-7 and HCT116 cell  
21 lines and 'mixed populations' of transfected cells  
22 were maintained in medium supplemented with 100µg/ml  
23 (MCF-7) or 1.5mg/ml (HCT116) G418 (from Life  
24 Technologies Inc).

25

26 **Generation of c-FLIP overexpressing cell lines.** c-  
27 FLIP<sub>L</sub> and c-FLIP<sub>S</sub> coding regions were PCR amplified  
28 and ligated into the pcDNA/V5-His TOPO vector  
29 according to the manufacturer's instructions (Life  
30 Technologies Inc.). HCT116p53<sup>+/+</sup> cells were co-  
31 transfected with 10µg of each c-FLIP expression  
32 construct and 1µg of a construct expressing a



1 puromycin resistance gene (pIRESpuro3, Clontech)  
2 using GeneJuice. Stably transfected HCT116 cells  
3 were selected and maintained in medium supplemented  
4 with 1µg/ml puromycin (Life Technologies Inc.).  
5 Stable overexpression of c-FLIP was assessed by  
6 Western blot analysis.

7  
8 **Western Blotting.** Western blots were performed as  
9 previously described (Longley et al., 2002). The  
10 Fas/CD95, Bcl-2 and BID (Santa Cruz Biotechnology,  
11 Santa Cruz, CA), caspase 8 (Oncogene Research  
12 Products, Darmstadt, Germany), PARP (Pharmingen, BD  
13 Biosciences, Oxford, England), c-FLIP (NF-6, Alexis,  
14 Bingham UK) DcR3 (Imgenex, San Diego, CA) mouse  
15 monoclonal antibodies were used in conjunction with  
16 a horseradish peroxidase (HRP)-conjugated sheep  
17 anti-mouse secondary antibody (Amersham, Little  
18 Chalfont, Buckinghamshire, England). FasL rabbit  
19 polyclonal antibody (Santa Cruz Biotechnology) was  
20 used in conjunction with an HRP-conjugated donkey  
21 anti-rabbit secondary antibody (Amersham). Equal  
22 loading was assessed using a β-tubulin mouse  
23 monoclonal primary antibody (Sigma).

24  
25 **Co-immunoprecipitation reactions.** 250µl of Protein A  
26 (IgG) or Protein L (IgM) Sepharose beads (Sigma) and  
27 1µg of the appropriate antibody were mixed at 4°C  
28 for 1 hour. Antibody-associated beads were washed  
29 three times with ELB buffer (250mM NaCl, 0.1%  
30 IPEGAL, 5mM EDTA, 0.5mM DTT, 50mM HEPES). Protein  
31 lysate (200-400µg) was then added, and the mixture  
32 rotated at 4°C for 1 hour. The beads were then

1 washed in ELB buffer five times and resuspended in  
2 100µl of Western sample buffer (250mM TRIS pH 6.8,  
3 4% SDS, 2% glycerol, 0.02% bromophenol blue)  
4 containing 10% β-mercaptoethanol. The samples were  
5 then heated at 95°C for 5 minutes and centrifuged  
6 (5mins/4,000rpm/4°C). The supernatant was collected  
7 and analysed by Western blotting.

8  
9 **Cell Viability Assays.** Cell viability was assessed  
10 by MTT (3-(4,5-dimethylthiazol-2-yl)-2,5-  
11 diphenyltetrazolium bromide, Sigma) assay (Mosmann,  
12 1983). To investigate drug-induced Fas-mediated  
13 apoptosis, cells were seeded at 2,000-5,000 cells  
14 per well on 96-well plates. After 24 hours, the  
15 cells were treated with a range of concentrations of  
16 5-FU, TDX, MTA or OXA for 24-72 hours followed by  
17 the agonistic Fas monoclonal antibody, CH-11 (MBL,  
18 Watertown, MA) for a further 24-48 hours. To assess  
19 chemotherapy/siRNA interactions, 20,000-50,000 cells  
20 were seeded per well on 24-well plates. Twenty-four  
21 hours later, the cells were transfected with FLIP-  
22 targeted (FT) or scrambled siRNA (SC). Four hours  
23 after transfection, the cells were treated with a  
24 range of concentrations of each drug for a further  
25 72-96 hours. MTT (0.5mg/ml) was added to each well  
26 and the cells were incubated at 37°C for a further 2  
27 hours. The culture medium was removed and formazan  
28 crystals reabsorbed in 200µl (96-well) or 1ml (24-  
29 well) DMSO. Cell viability was determined by reading  
30 the absorbance of each well at 570nm using a  
31 microplate reader (Molecular Devices, Wokingham,  
32 England).

1  
2 **Flow Cytometric Analysis.** Cells were seeded at  $1 \times 10^5$   
3 per well of a 6-well tissue culture plate. After 24  
4 hours, 5-FU, TDX or OXA was added to the medium and  
5 the cells cultured for a further 72 hours, after  
6 which time 250ng/ml CH-11 was added for 24 hours.  
7 DNA content of harvested cells was evaluated after  
8 propidium iodide staining of cells using the EPICS  
9 XL Flow Cytometer (Coulter, Miami, Fl).

10  
11 **siRNA transfections.** FLIP-targeted siRNA was  
12 designed using the Ambion siRNA target finder and  
13 design tool  
14 ([www.ambion.com/techlib/misc/siRNA\\_finder.html](http://www.ambion.com/techlib/misc/siRNA_finder.html)) to  
15 inhibit both splice variants of c-FLIP. Both c-FLIP-  
16 targeted (FT) and scrambled control (SC) siRNA were  
17 obtained from Xeragon (Germantown, MD). The FT siRNA  
18 sequence used was: AAG CAG TCT GTT CAA GGA GCA. The  
19 FL siRNA sequence used was: AAG GAA CAG CTT GGC GCT  
20 CAA. The control non-silencing siRNA sequence (SC)  
21 used was: AAT TCT CCG AAC GTG TCA CGT. siRNA  
22 transfections were performed on sub-confluent cells  
23 incubated in Optimem medium using the oligofectamine  
24 reagent (both from Life Technologies Inc) according  
25 to the manufacturer's instructions.

26  
27 **Statistical Analyses.** The nature of the interaction  
28 between the chemotherapeutic drugs and FLIP-targeted  
29 siRNAs was determined by calculating the combination  
30 index (CI) according to the method of Chou and  
31 Talalay (14). CI values were calculated from  
32 isobolograms generated using the CalcuSyn software

1 programme (Microsoft Windows). According to the  
2 definitions of Chou and Talalay, a CI value of 0.85-  
3 0.9 is slightly synergistic, 0.7-0.85 is moderately  
4 synergistic, 0.3-0.7 is synergistic and 0.1-0.3 is  
5 strongly synergistic. An unpaired two-tailed t test  
6 was used to determine the significance of changes in  
7 levels of apoptosis between different treatment  
8 groups.

9

## 10 RESULTS

11

12 **Example 1. c-FLIP<sub>L</sub> is up-regulated, processed and**  
13 **bound to Fas in response to 5-FU and TDX.**

14

15 Analysis of Fas expression in MCF-7 cells revealed  
16 that it was up-regulated by ~12-fold 72 hours after  
17 treatment with an IC60 dose 5-FU and was also highly  
18 up-regulated (by ~7-fold ) in response to treatment  
19 with an IC60 dose (25nM) of TDX (Fig. 1A). FasL  
20 expression was unaffected by each drug treatment,  
21 but appeared to be highly expressed in these cells.  
22 Expression of FADD was also unaffected by drug  
23 treatment. Somewhat surprisingly, neither caspase 8,  
24 nor its substrate BID were activated in 5-FU- or  
25 TDX-treated cells as indicated by a lack of down-  
26 regulation of the levels of procaspase 8 or full-  
27 length BID (Fig. 1A). Bcl-2 was highly down-  
28 regulated in response to each agent. Interestingly,  
29 c-FLIP<sub>L</sub> but not c-FLIP<sub>S</sub> was up-regulated by drug  
30 treatment. Furthermore, c-FLIP<sub>L</sub> was processed to its  
31 p43-form indicative of its recruitment and  
32 processing at the DISC (Fig. 1A). Expression of the

1 Fas decoy receptor DcR3 was unaltered by drug  
2 treatment in these cells.

3  
4 To further investigate the apparent inhibition of  
5 caspase 8 activation in 5-FU- and TDX-treated cells,  
6 we analysed the interaction between Fas and FasL  
7 following drug treatment. Co-immunoprecipitation  
8 reactions demonstrated that there was increased Fas-  
9 FasL binding following drug treatment (Fig. 1B),  
10 suggesting that the inhibition of caspase 8  
11 activation was occurring downstream of receptor  
12 ligation. In support of this, we found that drug  
13 treatment increased the interaction between Fas and  
14 p43- c-FLIP<sub>L</sub> (Fig. 1C). These results suggested the  
15 involvement of c-FLIP<sub>L</sub> in inhibiting drug-induced  
16 activation of Fas-mediated apoptosis in MCF-7 cells.

17  
18 **Example 2 Activation of drug-induced apoptosis by**  
19 **the Fas-targeted antibody CH-11 coincides with**  
20 **processing of c-FLIP<sub>L</sub>.** Expression of FasL by  
21 activated T cells and NK cells induces apoptosis of  
22 Fas expressing target cells in vivo. To mimic the  
23 effects of these immune effector cells in vitro, the  
24 agonistic Fas monoclonal antibody CH-11 was used.  
25 Cells were treated with either 5-FU or TDX for 72  
26 hours followed by 250ng/ml CH-11 treatment for 24  
27 hours. We found that CH-11 alone had little effect  
28 on apoptosis (Figs. 2A and B). Treatment with 5-FU  
29 alone for 96 hours resulted in a modest ~2-fold  
30 induction of apoptosis in response to 5µM 5-FU (Fig.  
31 2A). However, addition of CH-11 to 5-FU-treated  
32 cells resulted in a dramatic increase in apoptosis,

1 with a ~55% of cells in the sub-G1/G0 apoptotic  
2 phase following co-treatment with 5 $\mu$ M 5-FU and CH-  
3 11. Similarly, the combination of TDX with CH-11  
4 resulted in dramatic activation of apoptosis, with  
5 ~60% of cells in the sub-G1/G0 apoptotic phase  
6 following combined treatment with 25nM TDX and CH-11  
7 (Fig. 2B). We also examined the effect of CH-11 on  
8 apoptosis induced by the DNA-damaging agent OXA,  
9 which also potently induces Fas expression in MCF-7  
10 cells (Fig. 2C). Similar to its effect on 5-FU and  
11 TDX-treated cells, CH-11 induced apoptosis of OXA-  
12 treated cells, with ~50% of cells in the sub-G1/G0  
13 apoptotic phase (Fig. 2D).

14  
15 We subsequently analysed activation of the Fas  
16 pathway in MCF-7 cells following co-treatment with  
17 5-FU and CH-11. As already noted, treatment with 5-  
18 FU alone resulted in dramatic up-regulation of Fas,  
19 but had no effect on caspase 8 activation (Fig. 2E).  
20 However, co-treatment of MCF-7 cells with 5-FU and  
21 CH-11 resulted in a dramatic activation of caspase 8  
22 as indicated by complete loss of procaspase 8 (Fig.  
23 2E). Furthermore, cleavage of PARP (poly(ADP) ribose  
24 polymerase), a hallmark of apoptosis, was only  
25 observed in MCF-7 cells co-treated with 5-FU and CH-  
26 11 (Fig. 2E). We next analysed the kinetics of  
27 caspase 8 activation in 5-FU and CH-11 co-treated  
28 cells. Caspase 8 was potently activated 12 hours  
29 after addition of CH-11 to 5-FU pre-treated cells  
30 (Fig. 2F). Importantly, this coincided with complete  
31 processing of c-FLIP<sub>L</sub> to its p43-form (Fig. 2F). By  
32 24 hours after the addition of CH-11, neither

1 procaspase 8 nor c-FLIP<sub>L</sub> (both its full-length and  
2 truncated forms) was detected.

3  
4 Similarly, treatment of HCT116p53<sup>+/+</sup> cells with  
5 IC<sub>50</sub>(72h) doses of 5-FU (5μM) or oxaliplatin (1μM) for  
6 48 hours resulted in potent up-regulation of Fas  
7 expression (Fig. 8A), but only modest activation of  
8 caspase 8 and no PARP cleavage (Fig. 8B). However,  
9 co-treatment with each drug and CH-11 resulted in  
10 potent activation of caspase 8 and PARP cleavage  
11 (Fig. 8B). Activation of caspase 8 correlated with  
12 the complete processing of c-FLIP<sub>L</sub> to p43-FLIP<sub>L</sub> in  
13 drug and CH-11 co-treated cells (Fig. 8B).  
14 Furthermore, addition of CH-11 to 5-FU- and  
15 oxaliplatin-treated HCT116p53<sup>+/+</sup> cells resulted in  
16 ~4- and ~8-fold up-regulation of c-FLIP<sub>S</sub>  
17 respectively (Fig. 8B). These results suggested the  
18 involvement of c-FLIP in regulating Fas-mediated  
19 apoptosis in HCT116p53<sup>+/+</sup> cells following  
20 chemotherapy.

21  
22 We also examined the ability of CH-11 to activate  
23 apoptosis in the HCT116 colon cancer cell line. Fas  
24 was potently up-regulated in HCT116 cells 48 hours  
25 after treatment with 5-FU, TDX and OXA (Fig. 3A).  
26 Treatment with each drug alone or CH-11 alone for 48  
27 hours failed to significantly activate caspase 8 or  
28 induce PARP cleavage (Fig. 3B). However, treatment  
29 with each drug for 24 hours followed by CH-11 for a  
30 further 24 hours resulted in activation of caspase 8  
31 and PARP cleavage. Importantly, activation of  
32 caspase 8 correlated with processing of c-FLIP<sub>L</sub> in

1 drug and CH-11 co-treated cells (Fig. 3B).

2

3 To further test the hypothesis that the  
4 intracellular signal to commit to death receptor-  
5 mediated apoptosis in HCT116p53<sup>+/+</sup> cells following  
6 drug treatment was regulated by c-FLIP, the  
7 inventors generated HCT116p53<sup>+/+</sup> cell lines that  
8 overexpressed c-FLIP<sub>L</sub> or c-FLIP<sub>S</sub>. The HFL17 and HFL24  
9 cell lines both overexpressed c-FLIP<sub>L</sub> by ~6-fold  
10 compared to cells transfected with a LacZ-expressing  
11 construct (HLacZ), while the HFS19 and HFS44 cell  
12 lines overexpressed c-FLIP<sub>S</sub> by ~5- and ~10-fold  
13 respectively compared to the control cell line (Fig.  
14 9A). Growth inhibition studies (MTT assays) were  
15 carried out to determine the IC<sub>50(72h)</sub> dose for each  
16 chemotherapy in each cell line. It was found that  
17 overexpressing c-FLIP<sub>S</sub> had no significant effect on  
18 the IC<sub>50(72h)</sub> dose of any of the drugs, while c-FLIP<sub>L</sub>  
19 overexpression caused a moderate 1.7-2.0-fold  
20 increase in the IC<sub>50(72h)</sub> dose of oxaliplatin, but had  
21 no effect on the IC<sub>50(72h)</sub> doses of the other drugs  
22 (Table 1).

23

24 Flow cytometry revealed that c-FLIP<sub>L</sub> overexpression  
25 did not affect cell cycle arrest in response to  
26 chemotherapy, but had a marked effect on  
27 chemotherapy-induced apoptosis (Fig. 9B). For  
28 example, treatment with 5μM 5-FU for 72 hours  
29 resulted in cell cycle arrest at the G1/S phase  
30 boundary in each cell line, however the levels of  
31 apoptosis in the two c-FLIP<sub>L</sub>-overexpressing lines  
32 was significantly reduced compared to the control



1 cell line, with ~15% of HFL17 cells and ~17% of  
2 HFL24 cells in the sub-G<sub>1</sub>/G<sub>0</sub> apoptotic fraction  
3 compared to ~41% in the HLacZ cell line ( $p < 0.0001$ ,  
4 Fig. 9B). In contrast, the levels of apoptosis  
5 induced by 5-FU in the two c-FLIP<sub>S</sub>-overexpressing  
6 lines were actually somewhat higher than in the  
7 control HLacZ cell line. Similar results were  
8 obtained with the other drugs, as overexpression of  
9 c-FLIP<sub>L</sub> significantly decreased oxaliplatin- and  
10 CPT-11-induced apoptosis, whereas c-FLIP<sub>S</sub>  
11 overexpression failed to inhibit chemotherapy-  
12 induced apoptosis (Fig. 9B). The similar IC<sub>50</sub>(72h)  
13 doses observed in the c-FLIP<sub>L</sub>-overexpressing cell  
14 lines and the HLacZ cell line (Table 1) probably  
15 reflects the fact that c-FLIP<sub>L</sub> overexpression did  
16 not affect chemotherapy-induced cell cycle arrest,  
17 resulting in similar levels of growth inhibition  
18 despite the differences in drug-induced apoptosis  
19 observed in these cell lines.

20

21 **Example 4 Overexpression of c-FLIP<sub>L</sub> inhibits**  
22 **chemotherapy-induced Fas-mediated cell death.** To  
23 further investigate the role of c-FLIP<sub>L</sub> in  
24 regulating Fas-mediated apoptosis following drug  
25 treatment, we developed a panel of MCF-7 cell lines  
26 overexpressing c-FLIP<sub>L</sub>. We developed cell lines with  
27 5-10-fold increased c-FLIP<sub>L</sub> expression compared to  
28 cells transfected with empty vector (Fig. 4A). The  
29 c-FLIP<sub>L</sub> -overexpressing cell lines FL44 and FL64 and  
30 cells transfected with empty vector (EV68) were  
31 taken forward for further characterisation. Cell  
32 viability assays indicated that treatment of EV68

1 cells with 5-FU followed by CH-11 resulted in a  
2 highly synergistic decrease in cell viability  
3 (RI=2.06,  $p<0.0005$ ) (Fig. 4B). However, no  
4 synergistic decrease in cell viability was observed  
5 in 5-FU and CH-11 co-treated FL44 or FL64 cells,  
6 with RI values of 1.14 and 1.01 respectively (Fig.  
7 4B). Furthermore, 5-FU and CH-11 co-treatment  
8 resulted in caspase 8 activation and PARP cleavage  
9 in EV68 cells (Fig. 4C). In contrast, c-FLIP<sub>L</sub>  
10 overexpression in FL64 cells abrogated both  
11 activation of caspase 8 and PARP cleavage in  
12 response to 5-FU and CH-11 co-treatment (Fig. 4C).  
13  
14 We next examined the effect of c-FLIP<sub>L</sub>  
15 overexpression on Fas-mediated apoptosis following  
16 treatment with the antifolates TDX and MTA and the  
17 DNA-damaging agent OXA. All three drugs  
18 synergistically decreased cell viability in EV68  
19 cells when combined with CH-11 (Figs. 5A and B).  
20 However, this synergistic interaction was inhibited  
21 by c-FLIP<sub>L</sub> overexpression in both the FL44 and FL64  
22 cell lines (Figs. 5A and B). Analysis of caspase 8  
23 activation and PARP cleavage confirmed that Fas-  
24 mediated apoptosis in response to all three agents  
25 was attenuated by c-FLIP<sub>L</sub> overexpression. Combined  
26 treatment with each antifolate and CH-11 resulted in  
27 caspase 8 activation in EV68 cells, but not FL64  
28 cells (Fig. 5C). Similarly, PARP cleavage in  
29 response to the antifolates and CH-11 was inhibited  
30 in the FL64 cell line (Fig. 5C). Although some  
31 caspase 8 activation and PARP cleavage were observed  
32 in FL64 cells following co-treatment with 5 $\mu$ M OXA

1 and CH-11, this was much reduced compared to the  
2 EV68 cell line (Fig. 5D). These results indicate  
3 that c-FLIP<sub>L</sub> is a key regulator of Fas-mediated  
4 apoptosis in response to 5-FU, antifolates and  
5 oxaliplatin.

6  
7 Similar experiemnts were carried out using a number  
8 of other cell lines and chemotherapeutic agents in  
9 combination with CH-11. The results are shown in  
10 Figure 9C. Treatment with 50ng/mL CH-11 in the  
11 absence of chemotherapy induced a small degree of  
12 apoptosis in the HLacZ control cell line (data not  
13 shown). However, co-treatment with each chemotherapy  
14 and CH-11 resulted in high levels of apoptosis in  
15 the HLacZ cell line (Fig. 9C). High levels of  
16 apoptosis were also observed in the c-FLIP<sub>S</sub>-  
17 overexpressing cell lines HFS19 and HFS44 in  
18 response to chemotherapy and CH-11 (Fig. 9C). In  
19 contrast, c-FLIP<sub>L</sub> overexpression in the HFL17 and  
20 HFL24 cell lines dramatically inhibited apoptosis in  
21 response to co-treatment with each chemotherapy and  
22 CH-11 (Fig. 9C). So, overexpression of c-FLIP<sub>L</sub>, but  
23 not c-FLIP<sub>S</sub>, protected HCT116p53<sup>+/+</sup> cells from both  
24 chemotherapy-induced apoptosis and apoptosis induced  
25 in response to co-treatment with chemotherapy and  
26 the Fas agonist CH-11.

27  
28 **Example 6 siRNA-targeting of c-FLIP sensitises**  
29 **cancer cells to chemotherapy.**

30  
31 Having established that c-FLIP<sub>L</sub> overexpression  
32 protected MCF-7 and HCT116 cells from chemotherapy-

1 induced Fas-mediated cell death, we next designed a  
2 FLIP-targeted (FT) siRNA to inhibit both c-FLIP  
3 splice variants. Transfection with 10nM FT siRNA  
4 potently down-regulated expression of both c-FLIP  
5 splice variants in MCF-7 cells (Fig. 6A). Cell  
6 viability analysis of MCF-7 cells transfected with  
7 FT siRNA indicated that co-treatment with 5-FU  
8 resulted in a supra-additive decrease in cell  
9 viability (Fig. 6B,  $RI=1.56$ ,  $p<0.005$ ).  
10 Interestingly, transfection of MCF-7 cells with FT  
11 siRNA significantly decreased cell viability in the  
12 absence of co-treatment with 5-FU, with an  
13 approximate 50% decrease in cell viability in cells  
14 transfected with 2.5nM FT siRNA (Fig. 6B). A  
15 scrambled control (SC) siRNA that had no effect of  
16 FLIP expression, also had no effect on cell  
17 viability either alone or in combination with 5-FU  
18 (data not shown). The decrease in cell viability in  
19 response to FT siRNA alone appeared to be due to the  
20 induction of apoptosis, as transfection of FT siRNA  
21 in the absence of co-treatment with drug induced  
22 significant levels of PARP cleavage (Fig. 6C, lane  
23 2). Furthermore, combined treatment with FT siRNA  
24 and 5-FU resulted in potent cleavage of PARP (Fig.  
25 6C), indicating that the synergistic decrease in  
26 cell viability observed in MCF-7 cells co-treated  
27 with these agents was due to increased apoptosis.  
28  
29 FT siRNA also potently down-regulated FLIP<sub>L</sub> and FLIP<sub>S</sub>  
30 expression in HCT116 cells (Fig. 7A). Analysis of  
31 caspase 8 activation in siRNA-transfected HCT116  
32 cells indicated that FT siRNA alone (1nM) caused

1 some activation of caspase 8, as indicated by the  
2 decrease in the levels of p53/55 zymogen and  
3 appearance of the p41/43 cleavage products (Fig. 7B,  
4 lane 3). This was accompanied by some PARP cleavage.  
5 At higher concentrations ( $>5\text{nM}$ ), FT siRNA alone  
6 caused more potent activation of caspase 8 and PARP  
7 cleavage in HCT116 cells (Fig. 7C). Both 5-FU ( $5\mu\text{M}$ )  
8 and TDX ( $100\text{nM}$ ) caused some caspase 8 activation in  
9 mock and SC transfected HCT116 cells as indicated by  
10 the presence of p41/p43 caspase 8, although no PARP  
11 cleavage was observed in these cells (Fig. 7B). The  
12 most potent activation of caspase 8 was observed in  
13 cells co-treated with  $1\text{nM}$  FT siRNA and 5-FU or TDX,  
14 with decreased expression of the p53/55 zymogen and  
15 increased expression of both the p41/43 and p18  
16 caspase 8 cleavage products (Fig. 7B, lanes 6 and  
17 9). Furthermore, activation of caspase 8 in FT  
18 siRNA/chemotherapy-treated HCT116 cells was  
19 accompanied by potent PARP cleavage. Cell viability  
20 assays indicated that co-treatment with  $0.5\text{nM}$  FT  
21 siRNA and  $5\mu\text{M}$  5-FU resulted in a synergistic  
22 decrease in cell viability (Fig. 8A,  $\text{RI}=2.10$ ,  
23  $p<0.0005$ ). In contrast, SC siRNA had no significant  
24 effect on cell viability either in the presence or  
25 absence of 5-FU. Furthermore, co-treatment with FT  
26 siRNA and both TDX and OXA resulted in synergistic  
27 decreases in cell viability, with RI values of 1.68  
28 and 2.26 respectively (Figs. 8B and C). These  
29 results indicate that inhibition of c-FLIP  
30 expression in HCT116 and MCF-7 cells dramatically  
31 sensitised them to chemotherapy-induced apoptosis.  
32

1 Further evidence that siRNA-targeting of c-FLIP  
2 sensitises HCT116p53<sup>+/+</sup> cells to chemotherapy is  
3 shown in Figure 11. FLIP-targeted siRNAs were  
4 designed to down-regulate expression of both c-FLIP  
5 splice variants. Of several siRNAs tested, one FLIP-  
6 targeted (FT) siRNA potently down-regulated  
7 expression of both c-FLIP splice variants in  
8 HCT116p53<sup>+/+</sup> cells at nanomolar concentrations (Fig.  
9 11A). We used this FT siRNA to analyse the effect of  
10 down-regulating c-FLIP expression on drug-induced  
11 apoptosis. Interestingly, transfection with 0.5nM FT  
12 siRNA in the absence of chemotherapy induced  
13 significant levels of apoptosis (~26%) in  
14 HCT116p53<sup>+/+</sup> cells compared to cells transfected with  
15 control siRNA (~9%) as assessed by flow cytometric  
16 analysis of cells in the sub-G<sub>0</sub>/G<sub>1</sub> apoptotic fraction  
17 ( $p < 0.0001$ ; Fig. 11B). Importantly, co-treatment of  
18 FT siRNA transfected cells with an IC<sub>60</sub><sub>72h</sub> dose of 5-  
19 FU for 48 hours resulted in a supra-additive  
20 increase in apoptosis, with ~43% of cells undergoing  
21 apoptosis compared to ~11% in 5-FU-treated cells  
22 transfected with the control non-silencing siRNA  
23 ( $p = 0.0018$ ; Fig. 11B). The results following  
24 oxaliplatin treatment were even more dramatic, with  
25 ~61 % of cells co-treated with FT siRNA and  
26 oxaliplatin in the sub-G<sub>1</sub>/G<sub>0</sub> phase after 48 hours,  
27 compared to ~17% of cells co-treated with control  
28 siRNA and oxaliplatin ( $p < 0.0001$ ; Fig. 11B).  
29  
30 Analysis of caspase 8 activation in siRNA-  
31 transfected HCT116p53<sup>+/+</sup> cells indicated that 0.5nM  
32 FT siRNA alone caused some activation of caspase 8,

1 as indicated by the decrease in the levels of p53/55  
2 zymogen and appearance of the p41/43 cleavage  
3 products (Fig. 11C). Consistent with the cell cycle  
4 data, transfection with 0.5nM FT siRNA resulted in  
5 some PARP cleavage in the absence of chemotherapy.  
6 Treatment with 5µM 5-FU also caused modest caspase 8  
7 activation in mock-transfected cells and cells  
8 transfected with control siRNA (as indicated by the  
9 presence of p41/p43 caspase 8), however no PARP  
10 cleavage was observed in these cells (Fig. 11C). By  
11 far the most potent activation of caspase 8 was  
12 observed in cells co-treated with 0.5nM FT siRNA and  
13 5-FU, with decreased expression of the p53/55  
14 zymogen and increased expression of the p41/43  
15 caspase 8-cleavage product (Fig. 11C). Furthermore,  
16 activation of caspase 8 in FT siRNA/5-FU-treated  
17 HCT116p53<sup>+/+</sup> cells was accompanied by complete PARP  
18 cleavage. Similar results were obtained for the  
19 antifolate tomudex, which is a specific inhibitor of  
20 nucleotide synthetic enzyme thymidylate synthase (TS)  
21 (Fig. 11C). Furthermore, potent caspase 8 activation  
22 and PARP cleavage were observed in cells co-treated  
23 with FT siRNA and oxaliplatin after 24 hours,  
24 compared to cells treated with either agent  
25 individually (Fig. 11C). In light of these results,  
26 we also examined the effect of down-regulating c-  
27 FLIP on apoptosis induced by CPT-11, another  
28 chemotherapeutic drug currently used in the  
29 treatment of colorectal cancer. As with the other  
30 drugs, down-regulation of c-FLIP sensitised  
31 HCT116p53<sup>+/+</sup> cells to CPT-11-induced activation of  
32 caspase 8 and apoptosis (Fig. 10C).

1  
2 Given the more than additive effects of FT siRNA and  
3 chemotherapy on apoptosis in HCT116p53<sup>+/+</sup> cells, we  
4 carried out cell viability assays to determine  
5 whether the interactions were synergistic. Cell  
6 viability assays indicated that co-treatment with FT  
7 siRNA and 5-FU resulted in combination index (CI)  
8 values of <1 for 8/9 concentrations (Fig. 11D).  
9 According to the definitions of Chou and Talalay,  
10 the CI values for FT siRNA/5-FU co-treatment  
11 indicated that there was a moderate synergistic  
12 interaction for 4/9 concentration combinations  
13 examined and a synergistic interaction for a further  
14 4 concentrations (Fig. 11D). Co-treatment with FT  
15 siRNA and oxaliplatin resulted in synergistic  
16 decreases in cell viability for all concentrations  
17 examined, with CI values ranging from ~0.25-0.75  
18 (Fig. 3D). Similarly, combined treatment with CPT-11  
19 and FT siRNA resulted in synergistic or moderate  
20 synergistic decreases in cell viability with CI  
21 values ranging from ~0.50-0.85 (Fig. 11D). Control  
22 siRNA had no effect on cell viability in the  
23 presence or absence of any of the drugs (data not  
24 shown). Collectively, these results indicate that  
25 down-regulation of c-FLIP expression dramatically  
26 sensitises HCT116p53<sup>+/+</sup> cells to 5-FU-, oxaliplatin-  
27 and CPT-11-induced apoptosis.

28

29 **Example 7A The agonistic Fas monoclonal antibody CH-**  
30 **11 synergistically activates apoptosis in response**  
31 **to CPT-11 and TDX in a p53-independent manner**



1 The agonistic anti-Fas antibody CH-11 has been shown  
2 to activate the Fas/CD95 receptor and cause  
3 apoptosis. Lack of up-regulation of the Fas/CD95  
4 receptor in a p53 mutant colon cancer cell line  
5 abolished the synergistic interaction between 5-FU  
6 and CH-11. In our study treatment of the p53 wild-  
7 type and null cell lines with a range of each of the  
8 chemotherapy agents 5-FU, TDX, CPT-11 and  
9 Oxaliplatin followed 24 hours later by the addition  
10 of the anti-Fas antibody CH-11 (200ng/ml) for a  
11 further 48 hours resulted in significant synergy for  
12 all the drugs in the p53 wild-type setting, but in  
13 the p53 null cells this synergy was only seen with  
14 the topoisomerase-I inhibitor CPT-11 and the  
15 thymidylate synthase inhibitor TDX. There was no  
16 synergistic interaction seen at all with Oxaliplatin  
17 in the p53 null cells at any dose, and only slight  
18 interaction with 5-FU at the higher doses (Fig. 17).  
19 Propidium iodide staining of the HCT116 p53 wild-  
20 type and null cell lines treated with these  
21 chemotherapeutic agents for 24 hours followed by CH-  
22 11 50ng/ml for an additional 24 hours confirmed that  
23 a synergistic interaction is seen with each of the  
24 drugs and CH-11 in the p53 wild-type cells (Fig.  
25 18), whereas in the p53 null setting only treatment  
26 with CPT-11 and to a lesser extent with TDX resulted  
27 in significant synergy with CH-11 50ng/ml.

28

29

30 **Example 7B Effect of p53 inactivation on the synergy**  
31 **between CH-11 and 5-FU, CPT-11 and Oxaliplatin**

1     Activation of the Fas/CD95 receptor by its natural  
2     ligand FasL or the monoclonal antibody CH-11 results  
3     in the recruitment and activation of procaspase 8 at  
4     the DISC. Procaspase 8 is cleaved to its active  
5     subunits p41/43 and p18. Poly(ADP-ribose)polymerase  
6     (PARP) is normally involved in DNA repair and  
7     stability, and is cleaved by members of the caspase  
8     family during early apoptosis.  
9     Western blot analysis of the p53 wild-type and null  
10    cell lines treated with IC60 doses of these  
11    chemotherapeutic agents for 24 hours followed by a  
12    further 24 hours of the anti-Fas antibody CH-11  
13    (200ng/ml) resulted in PARP cleavage and activation  
14    of procaspase 8 (with generation of the active  
15    p41/43 and p18 subunits) in the p53 wild-type cell  
16    line for each drug (Fig. 19). In the p53 null cell  
17    line PARP cleavage and procaspase 8 activation  
18    following the addition of CH-11 was only seen  
19    following treatment with CPT-11.

20

21     **Example 7C Effect of p53 status on c-FLIP regulated**  
22     **chemosensitivity**

23

24     In order to determine whether down-regulation of c-  
25     FLIP would also sensitise p53 null HCT116 cells to  
26     chemotherapy-induced apoptosis, we transfected these  
27     cells with FT siRNA and co-treated them with  
28     chemotherapy (5-FU, oxaliplatin and CPT-11). The p53  
29     null cells (HCT116p53<sup>-/-</sup>) expressed higher levels of  
30     both c-FLIP splice forms than p53 wild type cells  
31     (Fig. 12A), but expression was effectively down-  
32     regulated by 1nM FT siRNA (Fig. 12B). Treatment of

1 the p53 null cells with 1nM FT siRNA alone resulted  
2 in a modest increase in apoptosis after 72 hours,  
3 with ~14% of cells in the sub-G<sub>0</sub>/G<sub>1</sub> fraction compared  
4 to ~9% in SC siRNA transfected cells (p=0.0081; Fig.  
5 12C). Co-treatment of FT siRNA-transfected cells  
6 with 5µM 5-FU significantly increased the apoptotic  
7 fraction to ~29% compared to ~14% of 5-FU/SC siRNA  
8 co-treated cells (p=0.0003; Fig. 12C). Treatment of  
9 FT siRNA-transfected HCT116 p53 null cells with 5µM  
10 oxaliplatin resulted in a highly significant  
11 increase in cells undergoing apoptosis compared to  
12 oxaliplatin/SC siRNA co-treated cells (~46% compared  
13 to ~27%, p<0.0001; Fig. 4C). FT siRNA also increased  
14 apoptosis of HCT116p53<sup>-/-</sup> cells in response to 1µM  
15 CPT-11 to ~33% compared to ~22% in SC/CPT-11 co-  
16 treated cells (p=0.0002; Fig. 12C). These results  
17 indicate that down-regulating c-FLIP expression  
18 significantly enhanced chemotherapy-induced  
19 apoptosis in p53 null HCT116 cells, in particular  
20 oxaliplatin-induced apoptosis.

21  
22 We further analysed the effect of down-regulating c-  
23 FLIP on the chemosensitivity of p53 null HCT116  
24 cells using the MTT cell viability assay. While  
25 greater than additive increases in apoptosis were  
26 detected for combined treatment with FT siRNA and 5-  
27 FU in HCT116p53<sup>-/-</sup> cells (Fig. 12C), cell viability  
28 assays identified slight synergy in only 2/9  
29 combinations (Fig. 12D). Similarly, the interaction  
30 between FT siRNA and CPT-11 was found to be  
31 moderately or slightly synergistic for only 3/9 drug  
32 combinations (Fig. 12D). So, although c-FLIP down-

1 regulation sensitised HCT116p53<sup>-/-</sup> cells to 5-FU-  
2 and CPT-11-induced apoptosis (Fig. 12C), cell  
3 viability assays indicated that fewer drug  
4 combinations were synergistic than in the p53 wild  
5 type parental cell line, and that the degree of  
6 synergy was less. However, co-treatment of  
7 HCT116p53<sup>-/-</sup> cells with oxaliplatin and FT siRNA was  
8 synergistic or moderately synergistic for all nine  
9 combinations analysed, with CI values ranging from  
10 ~0.35-0.85 (Fig. 12D), most likely reflecting the  
11 greater level of apoptosis induced for this  
12 combination than for the other chemotherapeutic  
13 drugs (Fig. 12C).

14

15 **Effect of c-FLIP on chemosensitivity in other**  
16 **colorectal cancer cell lines.** In order to determine  
17 whether c-FLIP is a general modulator of  
18 chemosensitivity in colorectal cancer, we extended  
19 these studies into two further colorectal cancer  
20 cell line models, namely the p53 wild type RKO cell  
21 line and the p53 mutant H630 cell line. Each cell  
22 line expressed both c-FLIP splice forms, and FT  
23 siRNA down-regulated c-FLIP protein in both lines  
24 (Fig. 13A). As in the HCT116 cell lines, down-  
25 regulation of c-FLIP sensitised both cell lines to  
26 apoptosis induced by 5-FU, oxaliplatin and CPT-11  
27 (Fig. 5B). In each case, the effect of co-treatment  
28 with chemotherapy and FT siRNA was more than  
29 additive. Of note, the sensitisation to CPT-11 was  
30 particularly marked in both lines, with ~43% of FT  
31 siRNA/CPT-11 co-treated RKO cells undergoing  
32 apoptosis compared to ~15% of SC siRNA/CPT-11 co-

1 treated RKO cells, and ~32% of FT siRNA/CPT-11 co-  
2 treated H630 cells undergoing apoptosis compared to  
3 ~12% of SC siRNA/CPT-11 co-treated H630 cells. MTT  
4 analyses indicated synergistic interactions between  
5 FT siRNA and each drug in RKO cells, with the  
6 majority of CI values below 0.75 for each drug (Fig.  
7 13C). The synergy was less pronounced in the H630  
8 cells, with the combination of FT siRNA and CPT-11  
9 being the most consistently synergistic or  
10 moderately synergistic (Fig. 13C).

11

12 Collectively, these results indicate that c-FLIP  
13 plays an important role in regulating chemotherapy-  
14 induced apoptosis in colorectal cancer cell lines.  
15 Furthermore, while both p53 wild type, mutant and  
16 null cell lines are sensitised to chemotherapy-  
17 induced apoptosis following down-regulation of c-  
18 FLIP, the extent of synergy would appear to be less  
19 in cell lines lacking functional p53.

20

21 **Potent knock-down of c-FLIP induces apoptosis in the**  
22 **absence of chemotherapy.** As already discussed,  
23 transfection of 0.5nM FT siRNA into HCT116p53<sup>+/+</sup>  
24 cells significantly increased apoptosis in the  
25 absence of co-treatment with chemotherapy (Fig.  
26 10B). When higher concentrations of FT siRNA were  
27 used to more completely knock down expression of c-  
28 FLIP in HCT116p53<sup>+/+</sup> cells, a dramatic decrease in  
29 cell viability (Fig. 14A) and a significant increase  
30 in PARP cleavage and apoptosis was observed (Fig.  
31 14B and C) in the absence of chemotherapy. A similar  
32 effect was observed in HCT116p53<sup>-/-</sup> cells, although

1 the extent of PARP cleavage and apoptosis was less  
2 than in the p53 wild type cell line (Fig. 14B and  
3 C). However, exposure of HCT116p53<sup>-/-</sup> cells to higher  
4 concentrations of FT siRNA for 72 hours resulted in  
5 levels of apoptosis that approached those observed  
6 in the p53 wild type parental cell line (Fig. 14D).  
7 The IC<sub>50(72h)</sub> doses of FT siRNA in the p53 wild type  
8 and null cell lines were ~0.7nM and ~2.5nM  
9 respectively as determined by MTT assay. FT siRNA  
10 also potently induced apoptosis in RKO and H630  
11 cells in the absence of chemotherapy (Fig. 14E and  
12 F). The IC<sub>50(72h)</sub> doses in these cell lines were  
13 calculated to be ~5nM in RKO cells and ~25nM in H630  
14 cells. These results indicate that c-FLIP may be a  
15 general determinant of colorectal cancer cell  
16 viability even in the absence of cytotoxic drugs.  
17 Furthermore, targeting c-FLIP induced apoptosis in  
18 p53 wild type, mutant and null and colorectal cancer  
19 cells, suggesting that it may represent an important  
20 new therapeutic target for treating this disease.

21  
22 Examination of the kinetics of c-FLIP down-  
23 regulation following FT siRNA transfection indicated  
24 that both splice forms were efficiently down-  
25 regulated as early as 8 hours post-transfection  
26 (Fig. 15A). This is in agreement with previous  
27 findings, which indicate that c-FLIP is rapidly  
28 turned over in cells following treatment with the  
29 protein synthesis inhibitor cycloheximide (16).  
30 Down-regulation of c-FLIP at 8 hours correlated with  
31 decreased levels of procaspase 8 and the onset of  
32 apoptosis as indicated by PARP cleavage (Fig. 15A).

1 This was more apparent for the higher concentration  
2 of FT siRNA (10nM). By 12 and 24 hours post-  
3 transfection, the p41/43-caspase 8 cleavage  
4 fragments could be detected in addition to the  
5 decrease in procaspase 8 levels and PARP cleavage in  
6 response to 1nM and 10nM FT siRNA (Fig. 15A). In  
7 agreement with the Western blot analysis, flow  
8 cytometry indicated that the onset of apoptosis  
9 following FT siRNA transfection occurred between 6  
10 and 12 hours (Fig. 15B). Therefore, c-FLIP down-  
11 regulation would appear to be tightly coupled to  
12 caspase 8 activation and the onset of apoptosis.

13

14 **Effect of specific targeting of c-FLIP<sub>L</sub> on**  
15 **apoptosis.** Our initial observation was that  
16 activation of apoptosis in chemotherapy/CH-11-  
17 treated HCT116p53<sup>+/+</sup> cells coincided with loss of  
18 full-length c-FLIP<sub>L</sub> (Fig. 9B). It was therefore  
19 possible that the effects on cell survival of down-  
20 regulating both c-FLIP splice variants were actually  
21 a result of the down-regulation of c-FLIP<sub>L</sub>. In  
22 addition, data from the c-FLIP overexpressing cell  
23 lines suggested that c-FLIP<sub>L</sub> was the more important  
24 regulator of chemoresistance (Fig. 10B). So, we  
25 designed an siRNA to specifically down-regulate the  
26 long splice form without affecting expression of c-  
27 FLIP<sub>S</sub> (Fig. 16A). Similar to the effect of the dual-  
28 targeted siRNA, specific down-regulation of c-FLIP<sub>L</sub>  
29 induced apoptosis of HCT116p53<sup>+/+</sup> cells in the  
30 absence of chemotherapy, as indicated by PARP  
31 cleavage (Fig. 8A) and flow cytometry (data not  
32 shown). Furthermore, combined treatment with FL

1 siRNA and each chemotherapy resulted in enhanced  
2 apoptosis (Fig. 16B) and highly synergistic  
3 decreases in cell viability (Fig. 16C). Similar  
4 synergistic decreases in cell viability were  
5 observed in the H630 and RKO cell lines (data not  
6 shown). These data suggest that down-regulation of  
7 c-FLIP<sub>L</sub> is sufficient to recapitulate the effects of  
8 down-regulating both splice variants and that, of  
9 the two splice forms, c-FLIP<sub>L</sub> may be the more  
10 critical regulator of colorectal cancer cell death.  
11  
12

### 13 DISCUSSION

14  
15 We found that the Fas death receptor was highly up-  
16 regulated in response to 5-FU, the TS-targeted  
17 antifolates TDX and MTA and the DNA-damaging agent  
18 OXA in MCF-7 breast cancer and HCT116 colon cancer  
19 cells, however, this did not result in significant  
20 activation of apoptosis. Expression of FasL by  
21 activated T cells and natural killer cells induces  
22 apoptosis of Fas expressing target cells in vivo  
23 (O'Connell et al., 1999). To mimic the effects of  
24 these immune effector cells in our in vitro model,  
25 we used the agonistic Fas monoclonal antibody CH-11.  
26 We found that CH-11 potently activated apoptosis of  
27 chemotherapy-treated cells, suggesting that the Fas  
28 signalling pathway is an important mediator of  
29 apoptosis in response to these agents in vivo. Many  
30 tumour cells overexpress FasL, and it has been  
31 postulated that tumour FasL induces apoptosis of  
32 Fas-sensitive immune effector cells, thereby



1 inhibiting the antitumor immune response (O'Connell  
2 et al., 1999). This hypothesis has been supported by  
3 both in vitro and in vivo studies (Bennett et al.,  
4 1998; O'Connell et al., 1997). The strategy of  
5 overexpressing FasL requires that the tumour cells  
6 develop resistance to Fas-mediated apoptosis to  
7 prevent autocrine and paracrine induction of tumour  
8 cell death. The lack of caspase 8 activation that we  
9 observed in response to chemotherapy suggests that  
10 Fas-mediated apoptosis may be inhibited in MCF-7 and  
11 HCT116 and cancer cells, but that co-treatment with  
12 CH-11 was sufficient to overcome this resistance and  
13 activate Fas-mediated apoptosis.

14  
15 Fas signalling may be inhibited by c-FLIP, which can  
16 inhibit caspase 8 recruitment to and activation at  
17 the Fas DISC (Krueger et al., 2001). Multiple c-FLIP  
18 splice variants have been reported, however, only  
19 two forms (c-FLIP<sub>L</sub> and c-FLIP<sub>S</sub>) have been detected at  
20 the protein level (Scaffidi et al., 1999). Both  
21 splice variants have death effector domains (DEDs),  
22 with which they bind to FADD, blocking access of  
23 procaspase 8 molecules to the DISC. c-FLIP<sub>L</sub> is  
24 processed at the DISC as it is a natural substrate  
25 for caspase 8, which cleaves it to generate a  
26 truncated form of approximately 43kDa (p43-FLIPL)  
27 (Niikura et al., 2002). Cleaved p43- c-FLIP<sub>L</sub> binds  
28 more tightly to the DISC than full-length c-FLIP<sub>L</sub>.  
29 c-FLIP<sub>S</sub> is not processed by caspase 8 at the DISC.  
30 c-FLIP<sub>L</sub> appears to be a more potent inhibitor of  
31 Fas-mediated cell death than c-FLIP<sub>S</sub> (Irmeler et al.,  
32 1997; Tschopp et al., 1998). Initially both pro-

1 apoptotic and anti-apoptotic effects were proposed  
2 for c-FLIP. However, enhanced cell death occurred  
3 mainly in experiments using transient over-  
4 expression and may have been due to excessive levels  
5 of these DED-containing proteins, which may have  
6 caused clustering of other DED-containing proteins  
7 including procaspase 8, resulting in caspase  
8 activation (Siegel et al., 1998). The data from cell  
9 lines stably over-expressing c-FLIP and from mice  
10 deficient in c-FLIP support an anti-apoptotic  
11 function for c-FLIP (Yeh et al., 2000).

12

13 We found that c-FLIP<sub>L</sub> was up-regulated and processed  
14 to its p43-form in MCF-7 cells following treatment  
15 with 5-FU and TDX. Furthermore, activation of  
16 caspase 8 and apoptosis in cells co-treated with  
17 chemotherapy and CH-11 coincided with processing of  
18 c-FLIP<sub>L</sub>. These results suggested that c-FLIP<sub>L</sub>  
19 regulated the onset of drug-induced Fas-mediated  
20 apoptosis in these cell lines. This hypothesis was  
21 further supported by data from overexpression and  
22 siRNA studies. c-FLIP overexpression abrogated the  
23 synergistic interaction between CH-11 and 5-FU, TDX,  
24 MTA and OXA by inhibiting caspase 8 activation.  
25 Furthermore, siRNA-targeting of both c-FLIP splice  
26 variants sensitised cells to these chemotherapeutic  
27 agents as determined by cell viability and PARP  
28 cleavage assays. Collectively, these results  
29 indicate that c-FLIP inhibits apoptosis in response  
30 to these drugs.

31

1 Surprisingly, we also found that siRNA-mediated  
2 down-regulation of c-FLIP<sub>L</sub> and c-FLIP<sub>S</sub> induced  
3 caspase 8 activation and PARP cleavage in the  
4 absence of co-treatment with chemotherapy (although  
5 co-treatment with drug enhanced the effect). The  
6 inventors found that overexpression of c-FLIP<sub>L</sub>  
7 protected HCT116 cells from chemotherapy-induced  
8 apoptosis and apoptosis induced following co-  
9 treatment with chemotherapy and the Fas agonistic  
10 antibody CH-11. In addition to blocking caspase 8  
11 activation, DISC-bound c-FLIP has been reported to  
12 promote activation of the ERK, PI3-kinase/Akt and  
13 NFκB signalling pathways (Kataoka et al., 2000;  
14 Panka et al., 2001). The NFκB, PI3K/Akt and ERK  
15 signal transduction pathways are associated with  
16 cell survival and/or proliferation, therefore, c-  
17 FLIP is capable of both blocking caspase 8  
18 activation and also recruiting adaptor proteins that  
19 can activate intrinsic survival and proliferation  
20 pathways (Shu et al., 1997). Furthermore, c-FLIP  
21 also inhibits procaspase 8 activation at the DISCs  
22 formed by the TRAIL receptors DR4 and DR5 (Krueger  
23 et al., 2001). rTRAIL induces apoptosis in a range  
24 of human cancer cell lines including colorectal and  
25 breast, indicating that the TRAIL receptors are  
26 widely expressed in tumour cells (Ashkenazi, 2002).  
27 It is possible that expression of DR4 and DR5 is  
28 tolerated in tumours because c-FLIP converts the  
29 apoptotic signal to one which promotes survival and  
30 proliferation. Thus, siRNA-mediated down-regulation  
31 of c-FLIP may induce apoptosis by inhibiting FLIP-

1 mediated activation of NF $\kappa$ B, PI3K/Akt and ERK and  
2 promoting activation of caspase 8 at TRAIL DISCs.

3

4 We have found that c-FLIP is a key regulator of Fas-  
5 mediated apoptosis in response to 5-FU, TS-targeted  
6 antifolates and OXA. Our results suggest that c-FLIP  
7 may be a clinically useful predictive marker of  
8 response to these agents and that c-FLIP is a  
9 therapeutically attractive target.

10

11 Furthermore, Our findings indicate that c-FLIP<sub>L</sub>  
12 overexpression inhibits apoptosis of colorectal  
13 cancer cells in response to the chemotherapeutic  
14 agents used in the treatment of colorectal cancer  
15 (5-FU, oxaliplatin and CPT-11). This has particular  
16 clinical relevance given the high incidence of c-  
17 FLIP<sub>L</sub> overexpression observed in colorectal cancer  
18 (6) and suggests that c-FLIP<sub>L</sub> overexpression may  
19 contribute to chemoresistance in colorectal cancer.  
20 Interestingly, c-FLIP<sub>S</sub> overexpression failed to  
21 protect colorectal cancer cells from chemotherapy-  
22 induced apoptosis, or apoptosis induced by co-  
23 treatment with chemotherapy and CH-11. These results  
24 would suggest that, of the two splice forms, c-FLIP<sub>L</sub>  
25 is the more important mediator of resistance to  
26 chemotherapy in colorectal cancer cells.

27

28 Our study indicates that down-regulating c-FLIP in a  
29 panel of colorectal cancer cells that have not been  
30 selected for drug resistance increases their  
31 sensitivity to a range of cytotoxic drugs with  
32 differing mechanisms of action. Furthermore, the

1 study has demonstrated that the down-regulation of  
2 c-FLIP alone can induce apoptosis .

3  
4 It would appear from our c-FLIP overexpressing cell  
5 lines and studies using a c-FLIP<sub>L</sub>-specific siRNA  
6 that the long splice form may be the more important  
7 in mediating survival of colorectal cancer cells,  
8 however conclusive proof of this will require the  
9 generation of a c-FLIP<sub>S</sub>-specific siRNA. The  
10 induction of apoptosis following c-FLIP knock-down  
11 is most likely mediated by death receptors such as  
12 Fas and DR5. We have previously shown that Fas is  
13 up-regulated in response to 5-FU in HCT116p53<sup>+/+</sup> and  
14 RKO cells, but not in HCT116p53<sup>-/-</sup> and H630 cells  
15 (39), while DR5 is constitutively expressed in both  
16 HCT116 cell lines and the RKO and H630 lines  
17 (unpublished observations). It is possible that  
18 knocking down c-FLIP expression (either in the  
19 presence or absence of chemotherapy) removes c-FLIP-  
20 mediated inhibition of caspase 8 activation at Fas  
21 and/or DR5 DISCs, leading to caspase 8-mediated  
22 activation of apoptosis. Indeed, our initial  
23 evidence suggests that the onset of apoptosis and  
24 caspase 8 activation following c-FLIP knock-down are  
25 tightly coupled. In addition to blocking caspase 8  
26 activation, DISC-bound c-FLIP has been reported to  
27 promote activation of the anti-apoptotic ERK, PI3-  
28 kinase/Akt and NF- $\kappa$ B signalling pathways (7, 8). So,  
29 it is also possible that loss of c-FLIP eliminates  
30 DISC-dependent up-regulation of these survival  
31 pathways, leading to enhanced susceptibility to  
32 apoptosis. In addition, a recent study has suggested

1 that c-FLIP<sub>L</sub> may have a non-DISC-dependent anti-  
2 apoptotic function by binding to and inhibiting pro-  
3 apoptotic signalling via p38 MAPK (40).

4  
5 The p53 tumour suppressor gene is mutated in 40-60%  
6 of colorectal cancers most often in the central DNA-  
7 binding core domain responsible for sequence-  
8 specific binding to transcriptional target genes  
9 (41). p53 has been reported to both  
10 transcriptionally up-regulate c-FLIP (42) and target  
11 it for ubiquitin-mediated degradation by the  
12 proteasome (43), suggesting that the effect of p53  
13 on c-FLIP expression is complex. In the present  
14 study, we consistently found that expression of both  
15 c-FLIP splice forms was higher in the p53 null  
16 HCT116 cell line compared to the isogenic p53 wild  
17 type line. We also examined how p53 status affected  
18 cell viability when c-FLIP was down-regulated.  
19 Although siRNA targeting of c-FLIP significantly  
20 enhanced chemotherapy-induced apoptosis in p53 null  
21 HCT116 cells, the effect was not as dramatic as in  
22 the p53 wild type line. Similarly, the induction of  
23 apoptosis after a 48 hour exposure to FLIP-targeted  
24 siRNA alone was greater in the p53 wild type  
25 setting. However, longer exposure times (72 hours)  
26 and higher concentrations (10-100nM) of FT siRNA  
27 induced levels of apoptosis in the HCT116 p53 null  
28 cell line that approached those observed in the p53  
29 wild type parental cell line. It is possible that  
30 the differential sensitivity of the p53 wild type  
31 and null cells to FT siRNA was at least partly due  
32 to the higher constitutive levels of c-FLIP

1 expression in the p53 null line. It may also reflect  
2 lower levels of basal and chemotherapy-induced  
3 expression of the p53-regulated genes encoding the  
4 Fas and DR5 death receptors in the p53 null cell  
5 line, which lowers its sensitivity to loss of c-FLIP  
6 expression. Of note, down-regulation of c-FLIP  
7 markedly enhanced apoptosis in response to  
8 oxaliplatin in the p53 null cells, which are usually  
9 highly resistant to oxaliplatin (15). Further  
10 analyses revealed that the effects of targeting c-  
11 FLIP on chemotherapy-induced apoptosis were not  
12 confined to the HCT116 lines, as similar results  
13 were obtained in the p53 wild type RKO and p53  
14 mutant H630 lines. Moreover, more potent knock down  
15 of c-FLIP with higher concentrations of siRNA  
16 triggered apoptosis in the absence of chemotherapy  
17 in both RKO and H630 cell lines. Collectively these  
18 results suggest that c-FLIP is an important  
19 regulator of cell survival in p53 wild type, null  
20 and mutant colorectal cancer cells in the presence  
21 and absence of chemotherapy.

22

23 These findings have direct clinical relevance as 5-  
24 FU/leucovorin/oxaliplatin (FOLFOX) and 5-  
25 FU/leucovorin/CPT-11 (FOLFIRI) combination  
26 chemotherapies are currently widely used in the  
27 treatment of advanced colorectal cancer, and FOLFOX  
28 has recently been demonstrated to improve 3-year  
29 survival compared to 5-FU/leucovorin in the adjuvant  
30 setting of the disease (78.2% versus 72.9%,  $p=0.002$ )  
31 (44). Furthermore, clinical studies have  
32 demonstrated significantly elevated c-FLIP

1 expression in colorectal and gastric tumours (6,  
2 45), suggesting that c-FLIP may not only be a  
3 relevant clinical target in colorectal cancer, but  
4 also in gastric cancer, where 5-FU-based  
5 chemotherapy regimens are also used. In conclusion,  
6 this study suggests that c-FLIP may represent an  
7 important clinical marker of drug resistance in  
8 colorectal cancer and that targeting c-FLIP, either  
9 alone, or in combination with standard  
10 chemotherapies has therapeutic potential for the  
11 treatment of this disease.

12



1  
2  
3 All documents referred to in this specification are  
4 herein incorporated by reference. Various  
5 modifications and variations to the described  
6 embodiments of the inventions will be apparent to  
7 those skilled in the art without departing from the  
8 scope and spirit of the invention. Although the  
9 invention has been described in connection with  
10 specific preferred embodiments, it should be  
11 understood that the invention as claimed should not  
12 be unduly limited to such specific embodiments.  
13 Indeed, various modifications of the described modes  
14 of carrying out the invention which are obvious to  
15 those skilled in the art are intended to be covered  
16 by the present invention.

17

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